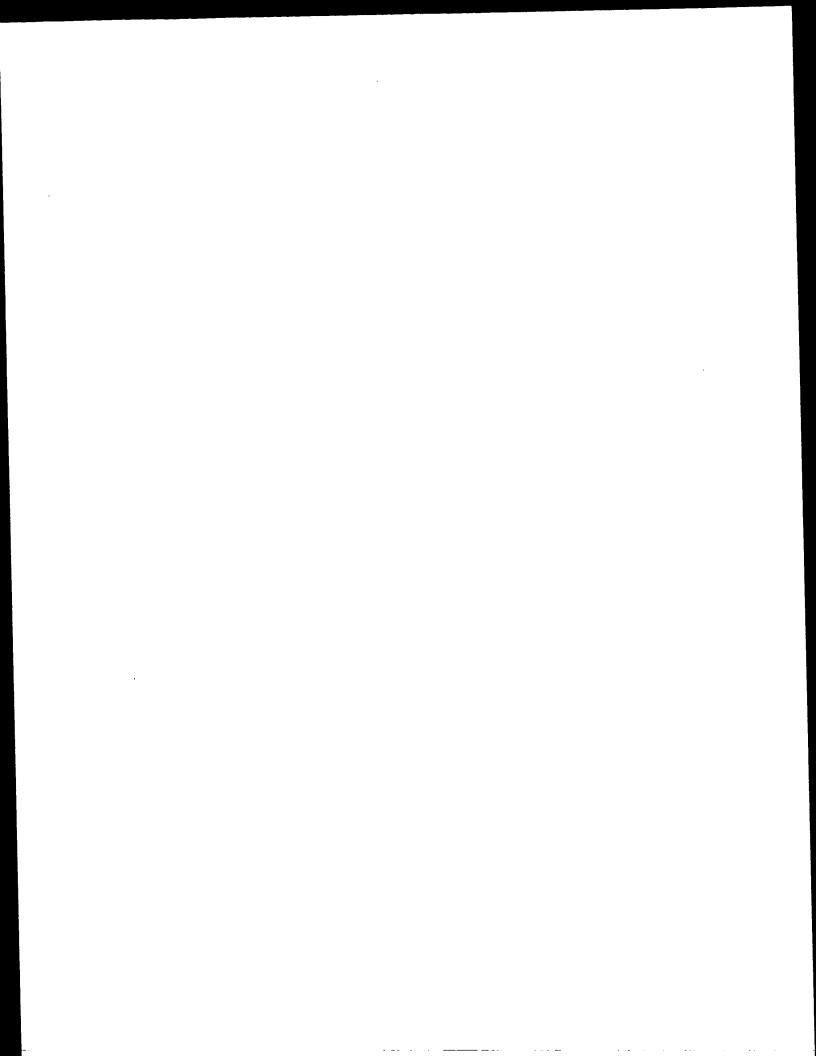
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PTO-1590 (9-90)



STIC-Biotech/ChemLib

From: Sent:

Schultz, James

To:

Monday, March 03, 2003 11:08 AM STIC-Biotech/ChemLib

Subject:

sequence search request for 10/001,844

Hello,

I need a length limited nucleotide sequence search performed on SEQ ID NO:3 (1576 nt long) in the above entitled case, where the maximum size of the returned hit is no longer than 50 nucleotides. Thank you very much,

Doug Schultz

J. Douglas Schultz, Ph.D AU 1635 (Biotechnology) Patent Examiner United States Patent and Trademark Office (703) 308-9355 (703) 746-3973 (fax) Office: CM1 12E18 Mail: CM1 11E12

Point of Contact: Beverly Shears Technical Info. Specialist CM1 1E05 Tel: 308-4994

Searcher:
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Sequence 9, Application US/09590211
GENERAL INFORMATION:
APPLICANT: Rouleau, Guy A.
APPLICANT: Brais, Bernard
TITLE OF INVENTION: Short GCG Expansions in the PAB II Gene
TITLE OF INVENTION: For Oculopharyngeal Muscular Dystrophy and Diagnostic
TITLE OF INVENTION: Thereof
FILE REFERENCE: 3028.1000-000
CURRENT FILING DATE: 2000-06-08
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
PRIOR FILING DATE: 1998-12-07
PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
LENGTH: 45
TYPE: DNA
ORGANISM: Homo Sapiens
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Best Local S
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      APPLICANT: Rouleau, Guy A.
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Local Similarity 83.7%;
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Sequence 25, Appl
Sequence 25, Appl
Sequence 25, Appl
Sequence 11755, A
Sequence 18755, A
Sequence 18755, A
Sequence 18755, A
Sequence 1563, Ap
Sequence 1563, Ap
Sequence 1563, Ap
Sequence 1563, Ap
Sequence 4569, Ap
Sequence 4569, Ap
Sequence 64136, A
Sequence 64136, A
Sequence 64136, A
Sequence 64137, A
                                                                                                                 45
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0;

APPLICANT:

Brais, Bernard

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US-09-590-211-8
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Sequence 8, Application US/09590211
GENERAL INFORMATION:
APPLICANT: Rouleau, Guy A.
APPLICANT: Brais, Bernard
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                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 16
LENGTH: SO
TYPE: DNA
                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                            Query Match
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/782,650 CURRENT FILING DATE: 2001-02-12 PRIOR APPLICATION NUMBER: US 09/324,079 PRIOR FILING DATE: 1999-06-01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/US00/14988
PRIOR FILING DATE: 2000-05-31
NUMBER OF SEQ ID NOS: 24
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PRIOR FILING DATE: 1999-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Edwards LifeSciences Corporat
TITLE OF INVENTION: Targeted Angiogenesis
FILE REFERENCE: 20553b-000611US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/590,211A
CURRENT FILING DATE: 2000-06-08
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR APPLICATION NUMBER: 2,218,199
PRIOR APPLICATION NUMBER: 2,218,199
                                                                                                                                                                                  1341 eceeceeceacacececceceeceececeecececec 1381
                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:annealed
                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                7 GCGGCGGCAGCGGCGGCGGCAGCGGCGGCGGCGGC 47
                                                                                                                                                                                                                                                      Local Similarity
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Falkner, Falko-Guenter
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                                                                                                                                                                                                                                  Conservative
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83.7%;
                                                                                                                                                                                                                                            2.0%; Score 31.4; DB 30; Length 50; 85.4%; Pred. No. 1.9e+05;
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PCT-US02-26129-14
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                                                                                                                                                                                            Sequence 14, Application PC/TUS0226129 GENERAL INFORMATION:
          CURRENT APPLICATION NUMBER: PCT/US02/26129
CURRENT FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 09/930,181
PRIOR FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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SOFTWARE: PatentIn version 3.1
                                                                                                                  TITLE OF INVENTION:
FILE REFERENCE: 16U
                                                                                                                                                         APPLICANT: OriGene Technologies APPLICANT: OriGene Technologies
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PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
PRIOR APPLICATION NUMBER: 2,218,199
PRIOR FILING DATE: 1997-12-09
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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CURRENT FILING DATE: 2000-06-08
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
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ORGANISM: Homo sapiens
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Similarity 82.5%;
                                                                                                                                                   OriGene Technologies
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                                                                                                              Serine Protein Kinase in Brain and Pancreas 101\ \text{PCT}
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 Mismatches

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; SEQ ID NO 14
; LENGTH: 50
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-195-072-14
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US-10-195-071-14
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US-10-195-072-14
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LENGTH: 50
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GENERAL INFORMATION:
APPLICANT: OriGene Technologies
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CURRENT FILING DATE: 2002-07-15
PRIOR APPLICATION NUMBER: US 09/930,181
PRIOR FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 18
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TITLE OF INVENTION: Full-Length Serine Protein Kinase in Brain and Pancreas
FILE REFERENCE: 16U 101 C1
                                                                    Matches
                                                                                                                                                                                                                                             TITLE OF INVENTION: Full-Length Serine Protein Kinase in Brain and Pancreas FILE REFERENCE: 16U 101 C2 CURRENT APPLICATION NUMBER: US/10/195,072 CURRENT FILING DATE: 2002-07-15 PRIOR APPLICATION NUMBER: US 09/930,181 PRIOR FILING DATE: 2001-08-16
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                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GCCCGGGCCGCGGCCTCGGCGCGCGGCGGCGGCGGCGGAAGC 49
                                                              / Match 1.8%; Score 28.2; DB 41;
Local Similarity 73.5%; Pred. No. 6.2e+05;
les 36; Conservative 0; Mismatches 13;
1 GGCGCGGGGCGCGCGCGGCGGCGGCGGCGGCGGCGGAAGC 49
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 Mismatches

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Pred. No. 6.2e+05;
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                                                                                                       Length 50;
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RESULT 9 US-09-590-211-7

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US-09-590-211-7
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                                                                                                 Sequence 289, Application US/09992665; GENERAL INFORMATION:
APPLICANT: Kaia Palm
                                                                                                                                                                      RESULT 11
US-09-992-665-289/c
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PRIOR FILING DATE: 1998-12-07
PRIOR APPLICATION NUMBER: 2,218,199
PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/590,211
CURRENT FILING DATE: 2000-06-08
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Best Local :
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APPLICANT: Brais, Bernard
TITLE OF INVENTION: SHORT GCG EXPANSIONS IN THE PAB II GENE
TITLE OF INVENTION: FOR OCCULOPHARYNGEAL MUSCULAR DYSTROPHY AND DIAGNOSTIC THEREFILE REFERENCE: 3028.1000-000
                                                                                                                                                                                                                                                                                                                                                 Matches
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CURRENT FILING DATE: 2000-06-08
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
PRIOR APPLICATION NUMBER: 2,218,199
PRIOR FILING DATE: 1997-12-09
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ORGANISM: Homo sapiens
                                    TITLE OF INVENTION: PROFILING TUMOR SPECIFIC MARKERS FOR THE TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF NEOPLASTIC DISEASE FILE REFERENCE: CEMINES.002A
CURRENT APPLICATION NUMBER: US/09/992,665
CURRENT FILING DATE: 2001-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                   1349 GACAGCGGCGGCGGGGACCGCGGGGGGGGGGGGGA 1385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 GGCGGCGGCGGCGGCGGCGGCGGCGCA 39
                                                                                                                                                                                                                                                                                                                                   1.7%;
Local Similarity 83.8%;
les 31; Conservation
                                                                                                                                                                                                                                                          3 GGCGGCGGCGGCGGCGGCGGCGGCGGCA 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 27.4; DB 22;
Pred. No. 8.4e+05;
                                                                                                                                                                                                                                                                                                                                                                     Score 27.4; DB 22;
Pred. No. 8.4e+05;
                                                                                                                                                                                                                                                                                                                                                     Mismatches
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US-09-590-211A-6
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APPLICANT: ROLLeau, Guy A.
APPLICANT: ROLLeau, Guy A.
APPLICANT: Brais, Bernard
TITLE OF INVENTION: SHORT GCG EXPANSIONS IN THE PAB II GENE
TITLE OF INVENTION: FOR OCULOPHARYNGEAL MUSCULAR DYSTROPHY AND DIAGNOSTIC THEREOF
FILE REFERENCE: 3028.1000-000
CURRENT TAPLICATION NUMBER: US/09/590,211A
CURRENT FILING DATE: 2000-06-08
PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
PRIOR APPLICATION NUMBER: Z,218,199
PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FRSTSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 2,218,199
PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/CA98/01133
PRIOR FILING DATE: 1998-12-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 3028.1000-000
CURRENT APPLICATION NUMBER: US/09/590,211
CURRENT FILING DATE: 2000-06-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Short GCG Expansions in the PAB II Gene TITLE OF INVENTION: for Oculopharyngeal Muscular Dystrophy and Diagnostic TITLE OF INVENTION: Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Rouleau, Guy A. APPLICANT: Brais, Bernard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 36
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          755 TCGGCCACGGTGCACCTGGAGCAGGGC 781
                                                                                                                                                                                                                                                                                                                                                                                                    4 GCGGCGGCGGCGGCGGCGGCGGCGGCA 36
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87.9%;
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 26.6; DB 22;
Pred. No. 1.1e+06;
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PCT-US00-19843-13
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                                                                                                                                                                 SOFTWARE: P
SEQ ID NO 13
LENGTH: 45
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   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 2
LENGTH: 36
                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-07-23
NUMBER OF SEQ ID NOS: 18
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                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: PCT/US00/19843
CURRENT FILING DATE: 2000-07-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                        OTHER INFORMATION: Description of Artificial Sequence:nucleic acid OTHER INFORMATION: linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cassandra L. Smith
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TARGETED
TITLE OF INVENTION: DNA DIFFERENTIAL DISPLAY
FILE REFERENCE: BU-042XX
CURRENT APPLICATION NUMBER: US/09/423,041A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: PT/US98/08616
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                                                                                                             FEATURE:
                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                 TYPE: DNA
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PRIOR FILING DATE: 1997-04-29
NUMBER OF SEQ ID NOS: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                   PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                   Herr, John C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                           00415-02
                                                                                                                                                                                                                                                                                                                                    Recombinant Antibody Directed Against Human Sperm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
   1.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
Score 26.4; DB 1;
Pred. No. 1.2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 26.4; DB 18;
Pred. No. 1.2e+06;
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               Length 45;
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US-10-031-783-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-031-783-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Pinus radiata US-09-606-755-17709
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SEQ ID NO 13
LENGTH: 45
                                                                          US-09-857-723-178/c
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Sequence 178, Application US/09857723 GENERAL INFORMATION:
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LENGTH: 50
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CURRENT FILING DATE: 2002-10-23
PRIOR APPLICATION NUMBER: 60/145,512
PRIOR FILING DATE: 1999-07-23
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APPLICANT: Norton, Elizabeth J.
APPLICANT: Deikman, Alan B.
TITLE OF INVENTION: Recombinant Antibody Directed Against Human Sperm
TITLE OF INVENTION: Antigen
FILE REFERENCE: 00415-02
FILE REFERENCE: 00415-02
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 17747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Polynucleotides, Materials Incorporating TITLE OF INVENTION: Them, and Methods for Using Them FILE REFERENCE: 11000.1041U6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Havukkala, Ilkka
APPLICANT: Shenk, Michael A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 17747
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                 19 GAGAGAGCGAGCGGGCGAGCCGGAGCGAAGGGGAAAGCGCAAGAGAGA 68
                                                                                                                                                                                                                                                                                                                                                                                                       Match 1.6%;
Local Similarity 70.0%;
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                                                                                                                                                                                                                           35;
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Pred. No. 1.2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                   Score 26; DB 23;
Pred. No. 1.4e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11;
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; NAME/KEY: satellite
; LOCATION: (1)...(50)
US-09-857-723-178
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US-09-857-723-199/c Sequence 199, Application US/09857723; GENERAL INFORMATION:
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                                                                                                                                                                                                                                          ; LOCATION: (1)...(50)
US-09-857-723-187
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NUMBER OF SEQ ID NOS: 292
SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: NUCLEIC ACID MOLECULES OF SHRIMP
FILE REFERENCE: 1322.1023-003
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TITLE OF INVENTION: NUCLEIC ACID MOLECULES OF SHRIMP
FILE REFERENCE: 1322.1023-003
                                                                                                                                                                                         Best
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PRIOR FILING DATE: 1998-12-10
NUMBER OF SEQ ID NOS: 292
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2001-06-07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Alcivar-Warren, Acacia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Litopenaeus vannamei
                                                                                                                                                                                                                                                                                            FEATURE:
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                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                        NAME/KEY: satellite
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                                                                                                                                    20 AGAGAGCGAGCGGGGGAGCGAGGAAGGGAAAGCGCAAGAGAGAGAG 69
                                                                                                                                                                                        Local Similarity
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Fan, Yongjun
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Fan, Yongjun
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pred. No. 1.4e+06;
0; Mismatches 15
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                                                                                                                                                                                           Score 26; DB 32;
Pred. No. 1.4e+06;
                                                                                                                                                                             Mismatches
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APPLICANT: Alcivar-Warren, Acacia

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; MOLECULE TYPE: other nucleic acid

; DESCRIPTION: /desc = "DNA"

US-08-715-713-17
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; NAME/KEY: satellite
; LOCATION: (1)...(50)
US-09-857-723-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17, Application US/08715713
GENERAL INFORMATION:
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                                                                                                                               TELEPHONE: (415) 705-8410
TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 17.
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CURRENT FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: 60/111,670
PRIOR FILING DATE: 1998-12-10
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TITLE OF INVENTION: NUCLEIC ACID MOLECULES OF SHRIMP
FILE REFERENCE: 1322.1023-003
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TYPE: DNA
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                       REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: UM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0:
FILING DATE: 19-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                           STRANDEDNESS:
                                                  TOPOLOGY:
                                                                                     TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 220 ECCITY: San Francisco
                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                    NAME:
                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE:
                                                                                                 LENGTH:
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                                                                                H: 50 base pairs
nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     220 Montomgery Street, Suite 2200
                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                             United States of America
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                                                                                                                                                                                                                                                                                                                                                                            Floppy disk
                                                              double
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                                                                                                                                ; ORGANISM: mus musculus US-09-396-196F-67174
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                                                                                                                                                                                SEQ ID NO 67174
LENGTH: 25
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                                                                     Matches
                                                                                                    Query Match
                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/396,196F
CURRENT FILING DATE: 2001-09-15
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FASTSEQ for Windows Version 4.0
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GENERAL INFORMATION:
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TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
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                                                                                                                                                                  TYPE: DNA
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
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TYPE: DNA
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nes 25; Conservative
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David Lockhart
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                                                                            1.6%; Score 25; DB 17;
100.0%; Pred. No. 2.1e+06;
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US-09-396-196F-67183

RESULT 24

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RESULT 25
US-09-396-196G-67173
; Sequence 67173, Application US/09396196G
; GENERAL INFORMATION:
; APPLICANT: Michael Mittmann
; APPLICANT: David Mack
; APPLICANT: David Tookhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: DNA; mus musculus; ORGANISM: mus musculus US-09-396-196F-67183
                                                                                                                                                                                                                                                                              ; ORGANISM: mus musculus US-09-396-196G-67173
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GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lookhart
APPLICANT: David Lookhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
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CURRENT FILING DATE: 2001-09-15
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 67183
LEBGTH: 25
                   Search completed: March 14, 2003, 01:02:17
Job time: 3791 secs
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PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 67173
LENGTH: 25
TYPE: DNA
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APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,1966
CURRENT FILING DATE: 1999-09-15
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975
       GENERAL INFORMATION:
APPLICANT: Feldmann, Richard J.; Global Determinants, Inc.
APPLICANT: Feldmann, Richard J.; Global Determinants, Inc.
TITLE OF INVENTION: Caulobacter crescentus complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT FILIGON NUMBER: US/10/287,787
CURRENT FILING DATE: 2003-03
NUMBER OF SEQ ID NOS: 27958
SOFTWARE: Proprietary
                                                                                                                                                                                                                                                                                                    SEQ ID NO 6
LENGTH: 30
                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               995
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987
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                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ANTISENSE MODULATION OF SHH EXPRESSION FILE REFERENCE: ISPH-0617
CURRENT APPLICATION NUMBER: US/10/001,844
CURRENT FILING DATE: 2001-11-16
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                                                                                                   Sequence 7843,
                                                                                                                                                                                                     Matches
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SEQ ID NO 7843
                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 49 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                          OTHER INFORMATION: PCR Probe
                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                              691 TATCCACTGCTCGGTGAAAGCAGAGAACTC 720
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US-10-287-787-14195

US-10-287-787-14822

US-10-287-787-16477
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US-60-427-836-652973

US-60-427-836-653150

US-60-427-836-654282

US-60-427-836-663449

US-10-367-892-21796
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US-10-314-578-1097

US-10-367-892-7821

US-10-299-054A-3456

US-10-299-054A-3456

US-10-287-787-13994

PCT-US02-24274-11

US-10-287-787-19712

US-10-287-787-10660

US-10-367-892-964
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US-10-299-054A-649
US-10-299-054A-4199
US-10-299-054A-10856
US-10-299-054A-10856
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Pred. No. 1.4e+C
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12674, A
14195, A
14822, A
16473, A
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654282,
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1097,
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13994, A
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; OTHER INFORMATION: Chromosome = 1 Strand = positive ConnectronObjectNumber US-10-287-787-7843
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TITLE OF INVENTION: Mycobacterium tuberculosis complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/299,054A
CURRENT FILING DATE: 2003-03-03
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                                                                                                                                                                                                                         SOFTWARE: Proprietary SEQ ID NO 10430 LENGTH: 43
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                                                              Matches
                                                                                           Query Match
                                                                                                                                                                                                                                                                              APPLICANT: Feldmann, Richard J.; GLOMAL DECEMBRICANT: Feldmann, Richard J.; GLOMAL DECEMBRICANT: FILE OF INVENTION: Mycobacterium tuberculosis complete genome. FILE REFERENCE: JIM Zegeer Law Offices - 703-684-8333 CURRENT APPLICATION NUMBER: US/10/299,054A CURRENT FILING DATE: 2003-03-03
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                                                                                                                                                                                      ORGANISM: Mycobacterium tuberculosis complete genome.
                                                                                                                                            LOCATION: (3933394)...(3933436)
OTHER INFORMATION: Chromosome = 1 Strand = positive
                                                                                                                                                                                                          TYPE: DNA
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75.0%;
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pred. No. 5.9e+04;
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Pred. No.
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                                                                                 Score 25.6; DB 7. Pred. No. 8.1e+04.
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US-10-299-054A-10450
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      Sequence 10450, Application US/10299054A
GENERAL IMPORMATION: Application US/10299054A
GENERAL IMPORMATION: Mycobacterium tuberculosis complete genome.
TITLE OF INVENTION: Mycobacterium tuberculosis complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/299,054A
CURRENT APPLICATION NUMBER: US/10/299,054A
UNUMBER OF SEQ ID NOS: 11910
SOPTWARE: Proprietary
SEQ ID NO 10450
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SOFTWARE: Proprietary
SEQ ID NO 10449
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Mycobacterium tuberculosis complete genome. FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333 CURRENT APPLICATION NUMBER: US/10/299,054A CURRENT FILING DATE: 2003-03-03
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TITLE OF INVENTION: Mycobacterium tuberculosis complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/01/299,054A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 11910
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LENGTH: 43
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OTHER INFORMATION: Chromosome = 1 Strand = positive ConnectronObjectNumber = 1252
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OTHER INFORMATION: Chromosome = 1 Strand = positive ConnectronobjectNumber = 12507
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Pred. No. 8
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US-10-303-778-10110/c Sequence 10110, Application US/10303778
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         CURRENT APPLICATION NUMBER: US/10/303,778
CURRENT FILING DATE: 2002-11-26
NUMBER OF SEQ ID NOS: 17608
SOFTWARE: Patentin version 3.1
SEQ ID NO 10110
LENGTH: 25
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                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: RosettaGenomics
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                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
                                                                                                                     FILE REFERENCE:
                                                                                                                            TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL TITLE OF INVENTION: REGULATORY GENES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/285,495
PRIOR FILING DATE: 2001-04-19
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/126,448 CURRENT FILING DATE: 2002-04-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: DISTITLE OF INVENTION: Synfile REFERENCE: CABI-004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Method for Diversifying the Chemical TITLE OF INVENTION: Composition of Proteins Produced in vivo by Genetically TITLE OF INVENTION: Disabling the Editing Function of their Aminoacyl tRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 50
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                             1353 GCGGCGGCGGGGACCGCGGGGGGGGGGGGGGG 1384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: synthetic oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: N = phosphorylated cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature LOCATION: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1339 CCGCGGCGGGACAGCGGCGGGGGGACCGCGGGGGGGGCGC 1378
                                                                                                                                                                                                                                                                                         45 GCGGCGGCGGCGGCGGCGGCGGCGGC 14
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schimmel, Pau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 de Crecy-Lagard, Valerie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hendrickson, Tamara L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nangle,
                                                                                                                                                                                                                                                                                                                                                                     Conservative
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87.5%;
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77.58;
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Pred. No. 8
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; ORGANISM: Homo sapiens US-10-303-778-10110
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                             US-60-427-836-454485
; Sequence 454485, Application US/60427836
; GENERAL INFORMATION:
; APPLICANT: Xue Mei Zhou
; TITLE OF INVENTION: Methods of Genetic Analysis of Rat
; FILE REFERENCE: 3527
; FILE REFERENCE: 3527
                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Mus musculus US-60-427-808-634065
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 296803
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 634065
LENGTH: 25
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TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Methods of Genetic Analysis of Mouse FILE REFERENCE: 3528 CURRENT APPLICATION NUMBER: US/60/427,808 CURRENT FILING DATE: 2002-11-20 NUMBER OF SEQ ID NOS: 982914
                                                                                                                                                                                                                                                           Matches
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ORGANISM: Mus musculus
 CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
CURRENT FILING DATE:
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                                                                                                                                                                                                                       1441 GGGCATCCACTGGTACTCGCAGCTG 1465
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Similarity 100.0%; Pred. No.
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100.0%; Pred. No.
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100.0%; Pred. No.
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 454485
RESULT 14
US-10-351-951-54
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                                                                                                                                                                                             US-09-912-293-16672
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                                                                                                                                                                                                                                                                                                          SEQ ID NO 16672
LENGTH: 50
                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/912,293
CURRENT FILING DATE: 2001-07-26
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                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 08/220,691
PRIOR FILING DATE: 1994-03-31
PRIOR APPLICATION NUMBER: 09/741,830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 08/103,744 PRIOR FILING DATE: 1993-08-09
                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/813,155 PRIOR FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (50)..(50)
OTHER INFORMATION: n is equal to a,t,g, or c
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                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                             5 GGCAGCCAGCGAGGGAGAGAGCGAGCGAGCCAGCGAGGAAGGG 53
                                                               APPLICATION NUMBER: 08/221,623 FILING DATE: 1994-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 2001-05-18
APPLICATION NUMBER: 08/196,362
FILING DATE: 1994-02-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 09/859,490
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APPLICATION NUMBER: 08/
FILING DATE: 1993-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 1994-02-15
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                                                                                                                                Conservative
                                                                                                                                                 1.6%;
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pred. No.
                                                                                                                                     0
                                                                                                                                                 Score 25; DB 6;
Pred. No. 1e+05;
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                                                                                                                                     Mismatches
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                                                                                                                                                                     DB 6;
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GENERAL INFORMATION:
APPLICANT: Stefanson, Stefan E.
TITLE OF INVENTION: GENE LINKED TO OSTEOARTHRITIS

Application US/10351951

Sequence 54,

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; TYPE: DNA
; ORGANISM: Mus musculus
US-60-427-808-121712
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Sequence 121712, Application US/60427808

GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                   RESULT 16
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; ORGANISM: Homo sapiens
US-10-351-951-54
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                                                                                                                 CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 121712
LENGTH: 25
  Best Local Similarity
                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                     APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1 SEQ ID NO 43
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APPLICANT: Ingham et al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 54
LENGTH: 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 08/674509
PRIOR FILING DATE: 1996-07-07
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/711,724A
CURRENT FILING DATE: 2000-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SCREENING ASSAYS FOR HEDGEHOG AGONISTS AND ANTAGONISTS FILE REFERENCE: HMSU-P14-006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/351,951
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: 10/057,312
PRIOR FILING DATE: 2002-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/431,538 PRIOR FILING DATE: 2002-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2345.2043-004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1310 CTGGCTGCACTGGCGCCCCGCGCACGGACCGCGGCGGGACAGCGGCGG 1359
                                                                                                                                                                                                                                                                                                                                                                                                                                       524 ACCGAGGGCTGGGACGAAGATGGC 547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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Local Similarity 68.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24;
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1.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.5%; Score 24; DB 6; Length 24; 100.0%; Pred. No. 1.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Pred. No. 1.8e+05;
                Score 23.4;
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                DB 9;
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              Length 25;
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RESULT 19
US-60-427-808-765746
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US-60-427-808-634064
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           CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe Sequence Listing SEQ ID NO 765746
LENGTH: 25
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                                                                                                                                                                                 Sequence 765746, Application US/60427808 GENERAL INFORMATION:
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SEQ ID NO 634064
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 634064, Application US/60427808 GENERAL INFORMATION:
                                                                                                                   APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
COGNUMBER OF SEQ ID NOS: 982914
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TYPE: DNA
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                              1441 GGGCATCCACTGGTACTCGCAGCTG 1465
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TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                       96.0%;
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                                           Sequence Listing Generator V 1.1

 Mismatches

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Pred. No. 1.
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Pred. No. 1.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     .8e+05;
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; ORGANISM: Mus musculus 
US-60-427-808-765746
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US-60-427-808-807373
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                                                                                                                                                                                                                                                                                                                ; ORGANISM: Mus musculus US-60-427-808-879497
                                                                                                                   US-60-427-836-90500
                                                                                                                                  RESULT 22
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LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 879497
LENGTH: 25
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3528
                                                                                Sequence 90500, Application US/60427836 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe Sequence Listing Generator
                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                 Best Local
                                                                                                                                                                                                                                                                             Query Match
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CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527
CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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                                                                                                                                                                                                   876 TGGACCGCGACGACGGCGCCAAGAA 900
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Local Similarity 96.0%;
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Similarity 96.0%;
24; Conservative
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Pred. No. 1.8e+05;
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                                                                                                                                                                                                                                                   0;
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Pred. No. 1
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Pred. No. 1.8e+05;
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                                                                                                                                                                                                                                                                                      Length 25;
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; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-60-427-836-90500
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US-60-427-836-238304
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                                                                                                                                                                                                                              ; ORGANISM: Rattus norvegicus US-60-427-836-454484
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RESULT 25
US-60-427-836-5917/91
; Sequence 5917/91, App.
; GENERAL INFORMATION:
                                                                                                   Вb
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CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 238304
LENCTH: 25
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1;
SEQ ID NO 9500
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Best Local Similarity
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                                                                                                                                                                                                                                                              SEQ ID NO 454484
LENGTH: 25
TYPE: DNA
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TITLE OF INVENTION: Methods of Genetic Analysis of Rat
                                                                                                                                                                  Matches
                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 699466
                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Methods of Genetic Analysis of Rat FILE REFERENCE: 3527
                                                                                                                                                                                                                                                                                                           SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xue Mei Zhou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Rattus norvegicus
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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I15914 Sequence 15

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AR135731 Sequence
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                                                                                                                   H.sapiens m
Synthetic I
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                                                                                     Sequence 18
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AX146613/c
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AX146614/c
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1. (bases 1 to 50)
                                                                                                                                                                                                                                                                                                                      49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BIOGEN, INC. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hedgehog fusion proteins and uses Patent: WO 0134654-A 44 17-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Strauch, K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AX146613 50 bp
Sequence 75 from Patent W00134654.
AX146613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
                                                             Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 50)
                                                                                                                                                                50 bp
Sequence 76 from Patent W00134654.
AX146614
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           Hedgehog fusion proteins and uses Patent: WO 0134654-A 76 17-MAY-2001; BIOGEN, INC. (US)
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                                                     Strauch, K
                                                                                                                          human
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/db_xref="taxon:9606"
11 c 15 g 1
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/db_xref="taxon:9606"
17 c 13 g 1
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Pred. No. 1.
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Local Similarity 94.0%;
nes 47; Conservating
       45;
                                                                                                                     BIOGEN, INC.
                                                                                                                            Hedgehog fusion proteins and uses Patent: WO 0134654-A 70 17-MAY-2001;
                                                                                                                                                       Strauch, K.
                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                Sequence 70 from Patent W00134654. AX146608
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                                                                                                                                                                                                              human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BIOGEN, INC. (US)
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      Conservative
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                                                             /db_xref="taxon:9606"
13 c 18 g
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                                                                                /organism="Homo sapiens"
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/db_xref="taxon:9606"
               2.8%;
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          Score 43.8; DB 6;
Pred. No. 3.8e+04;
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Pred. No. 2.5e+04;
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Pred No. 1.1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   44;
                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                   BIOGEN, INC. (US)
                                                                                                                                                                                                                        Hedgehog fusion proteins and uses Patent: WO 0134654-A 74 17-MAY-2001;
                                                                                                                                                                                                                                                      Strauch, K.
                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                 Sequence 74 from Patent W00134654.
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Sequence 71 from Patent W00134654.
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/db_xref="taxon:9606"
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97.8%;
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Pred. No. 3.8e+04;
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DEFINITION

Sequence 77 from Patent W00134654

DNA

linear

PAT 31-MAY-2001

LOCUS

AX146615

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ACCESSION
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AX146617
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AX146615.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strauch, K
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                      Sequence 73 from Patent W00134654.
AX146611
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  Strauch, K.
                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                    Homo sapiens
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Pred. No. 6.1e+04;
0; Mismatches 5
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                   Edwards Lifesciences Corporation (US) (AT)
                                                                                               Targeted angiogenesis
Patent: WO 0075329-A 16 14-DEC-2000;
                                                                                                                                     Levine, A.J., Mitterer, A., Falkner, F.G., Scheiflinger, F. and
                                                                                                                                                                               synthetic construct
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1 (bases 1 to 50)
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/db_xref="taxon:9606"
16 c 17 g
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                /organism="synthetic construct"
/db_xref="taxon:32630"
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Pred. No. 9.2e+04;
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                                  oligonucleotide
Patent: US 6319672-A 35 20-NOV-2001;
                                                    l (bases 1 to 50)
Crouzet,J., Scherman,D., Wils,P., Blanche,F. and Cameron,B.
Purification of a triple helix formation with an immobilize
                                                                                                                                                              Sequence 35 from patent US AR178318
                                                                                                               Unknown
                                                                                                                            Unknown.
                                                                                                                                                      AR178318.1
                                                                                                                                                                                          AR178318
                                                                                                                                                                                                                                                                                                        . Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (04-OCT-1999) Surgery, Cedars Sinal Medical Center, 8700 Beverly Blvd., Los Angeles, CA 90048, US Related entry: NM_000193
                                                                                                   Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Palm,K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Palm,K., Salin-Nordstrom,T., Levesque,M.F. and Neuman,T. Fetal and adult human CNS stem cells have similar molecular characteristics and developmental potential Brain Res. Mol. Brain Res. 78 (1-2), 192-195 (2000)
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AJ270316.1 GI:9557893
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                                                                                                                                                                                                                                                                                                                                                               /note="PCR antisense primer for sonic hedgehog (Drosophila) homolog (SHH)" 11 \ c \qquad 9 \ g \qquad 4 \ t
                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
1. .27
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CGGCGGCGGCGGCGGCGGCGGCGGCGG 33
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Schalling,M., Hudson,T.J. and Housman,D.E.
Direct detection of expanded nucleotide re
Patent: US 5695933-A 7 09-DEC-1997;
                                                                                                                                                                                                                                                                                 Sequence
184406
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11 c 22 q
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                                                                                                                                                                      repeats in the human genome
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                                                                                                                                                        1 (bases 1 to 50)
Drazen,J.M., In,K.-H., Asano,K., Beier,D. and Grobholz,J.
5-Lipoxygenase gene polymorphisms and their use in classifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6 from patent US 6355434.
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AR063105
AR063105.1 GI:5990796
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Location/Qualifiers
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Sequence 43 from patent US 6165747.
AR122632
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Ingham, P.W., McMahon, A.P. and Tabin, C.J.
                                                                                                                                                                                                                                                                                                                                                                        Unclassified.
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(bases 1 to 24)
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Pred. No. 7.4e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 24)
Ingham, P.W., McMahon, A.P. and Tabin, C.J.
Nucleic acids encoding hedgehog proteins
Patent: US 6271363-A 43 07-AUG-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids encoding hedgehog proteins Patent: US 6165747-A 43 26-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 43 from patent US 6271363. AR164260
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   24;
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AR208932
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                                                                                                             vertebrate embryonic pattern-inducing proteins
patent: US 6384192-A 43 07-MAY-2002;
                                                                                                                                                                                                             AR208932.1
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                                                                                                                                                             Unclassified.
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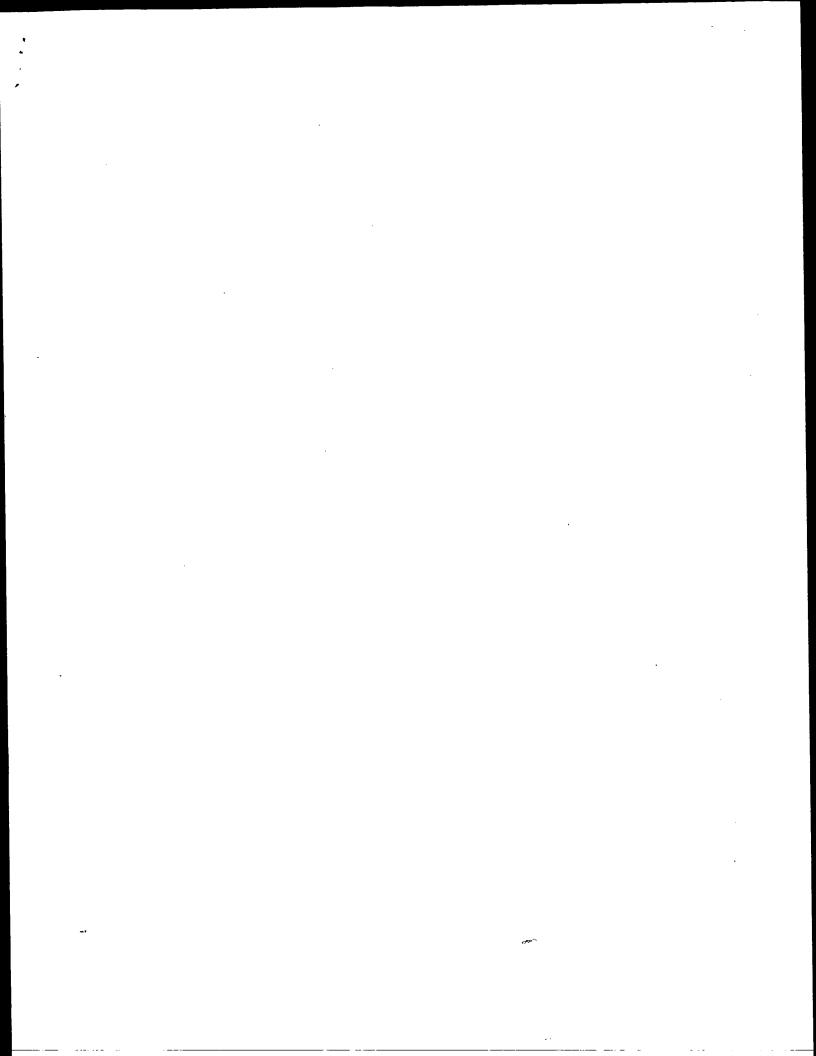
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synthetic construct. synthetic construct artificial sequences
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                                                                      Malaria parasitic epitope (T-cell).
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Schalling,M., Hudson,T.J. and Housman,D.E.
Direct detection of expanded nucleotide repeats in
Patent: US 5695933-A 2 09-DEC-1997;
                                                  A01419.1
                                                              A01419
                                                                                     A01419
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Huang, G.P., Rhode, P.R., Stinson, J.R. and Wong, H.C.
Method involving display of protein binding pairs on the surface of bacterial pili and bacteriophage
Patent: US 5516637-A 13 14-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                 I84401.1
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Pred. No. 1.1e+07;
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Pred. No. 8.9e+06;
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Job time: 4408 secs

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Search completed: March 13,
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Local Similarity 86.7%;
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1 (bases 1 to 32)
Oerum, H. and Seeger, C.
METHOD FOR GENERATING MULTIPLE DOUBLE STRANDED NUCLEIC ACIDS PATENT: WO 9720068-A 5 05-JUN-1997;
BOEHRINGER MANNHEIM GMBH (DE)
Location/Qualifiers
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                                                                                                                                                                                                                                                                             unidentified
                                                                                                                                                                                                                                                                                                                            Sequence 5 from Patent WO9720068.
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/db_xref="taxon:32644"
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/db_xref="taxon:32630"
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Pred. No. 1.5e+07;
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Maximum Match
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AAZ68973 ABAF30323 ABAF30323 ABAF30323 ABAF30323 ABAF30323 ABAF30323 ABAF30323 AAG02970 AAD16404 AAA35091 AAA71296 AAA735091 AAA741693 AAAC128563 AAAC29415 AAL29649 AAL29649 AAL29649 AAL29649 AAL29649 AAL29649 AAL2967 AAL31322 AAL31322 AAL31322 AAL34169 AAL3454331 AAL34547 AAL34547 AAL34547 AAL34547 AAL34547 AAL34547 AAL34547 AAL34547 AAL34547 AAA77627 AAAS15627 AAA77627 AAAS15778 AAA799891 AAA799891 AAA77627 AAAS13789 AAA34589 AAA33467 AAAS13789 AAA33467 AAAS13789 AAA33467 AAAS33643 AAA776227 AAAS33643 AAAT76227 AAAS33643 AAAA34589 AAAA333467 AAA2448716	900 17.4 1.1 1.2 903 17.4 1.1 1.1 1.2 906 17.4 1.1 1.1 1.2 908 17.4 1.1 1.1 1.2 908 17.4 1.1 1.1 1.2 908 17.4 1.1 1.2 908 17.4 1.1 1.2 908 17.4 1.1 1.2 918 17.4 1.1 1.2 918 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.4 1.1 1.2 928 17.2 1.1 1.2 938 17.2 1.2 1.2 1.2 1.2 1.2 1.2 1.2 1.2 1.2 1	17.4 1.1 17.4 1.1
Human map-related Human gamma 1 cons DNA binding molecu Chicken tenascin the Gene delivery fusi Human oligo 8 to c PCR primer #11. U PIV-1 zinc binding BMP-3 5' end probe Oligonucleotide who Oligonucleotide pro Human SNP oligonuc Human SIlent SNP c Human C/EBP-beta antisen Human C/EBP polynulatory Immunostimulatory Imm	900 17.4 1.1 500 2.916 17.4 1.1 500 2.906 17.4 1.1 500 2.907 17.4 1.1 500 2.908 17.4 1.1 500 2.908 17.4 1.1 500 2.916 17.4 1.1 500 2.916 17.4 1.1 500 2.916 17.4 1.1 500 2.916 17.4 1.1 500 2.916 17.4 1.1 500 2.916 17.4 1.1 500 2.927 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.4 1.1 500 2.928 17.2 1.1	17.4 1.1 47 17.4 1.1 47 17.4 1.1 47 17.4 1.1 47 17.4 1.1 48 17.4 1.1 48 17.4 1.1 48 17.4 1.1 50 17.4 1.1 50 17.4 1.1 50
Human map-related Human gamma 1 cons DNA binding molecu Chicken tenascin t Gene delivery fusi Human oligo 8 to c PCR primer #11. HIV-1 zinc binding BMP-3 5' end probe Oligonucleotide co Oligonucleotide ha Human secreted pro Human secreted pro Human swp oligonuc Human SNP oligon	900 17.4 1.1 50 22 903 17.4 1.1 50 22 906 17.4 1.1 50 22 908 17.4 1.1 50 22 908 17.4 1.1 50 22 908 17.4 1.1 50 22 908 17.4 1.1 50 22 908 17.4 1.1 50 22 908 17.4 1.1 50 22 910 17.4 1.1 50 22 911 17.4 1.1 50 22 912 17.4 1.1 50 22 913 17.4 1.1 50 22 915 17.4 1.1 50 22 92 92 92 92 92 92 92 92 92 92 92 92	17.4 1.1 47 2: 17.4 1.1 47 2: 17.4 1.1 47 2: 17.4 1.1 47 2: 17.4 1.1 48 1: 17.4 1.1 48 2: 17.4 1.1 48 2: 17.4 1.1 48 2: 17.4 1.1 49 2: 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 17 17.4 1.1 50 21 17.4 1.1 50 21 17.4 1.1 50 21
	900 17.4 1.1 50 21 AACLESSS 302 17.4 1.1 50 22 AACLESSS 309 17.2 1.1 2.1 2.1 2.1 2.1 2.1 2.1 2.1 2.1 2	17.4 1.1 47 21 AA268973 17.4 1.1 47 22 AAF30323 17.4 1.1 47 24 ABK82504 17.4 1.1 47 24 ABA99916 17.4 1.1 47 24 ABA99916 17.4 1.1 48 16 AAT02970 17.4 1.1 48 22 AA56404 17.4 1.1 48 22 AA513008 17.4 1.1 49 22 AA513008 17.4 1.1 50 17 AA35091 17.4 1.1 50 17 AA737108 17.4 1.1 50 21 AAA77273 17.4 1.1 50 21 AAA77273

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AAZ58685
AAZ44227
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                                                                                                                                                                                                  AAT38703
                                                                                                                                                  AAQ1
                                                                                                                                                                  AAQ51915
                                                                                                                                                                       AAX59829
                                                                                                                                                                                       AAQ14197
                                                                             AAV5
                                 AAT14495
                                                             ABL4
                                                                       AAV50609
           AAX36430
                                                                                                                                                             18455
                                                                                                                                                                                                                        Synthetic PCR prim
Anti-CD4 antibody
                                                                                            Human glucagon rec
Vector pVAC1 const
                                                                                                                                             Sense oligonucleot
                                                                                                                                                        Mutant
                                                                                                                                                                  c-myc mRNA ribozym
                                                                                                                                                                        Probe
                                                                                                                                                                             Probe used
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                                                                                                                                                                                         Probe P1446 for
                                                                                                                                                                                             PCR primer
                                                                                                                                                                                                        Permutein linker e
                                                                                                                                                                                                               806.077 scFv antib
                                                            Maize polymorphic Probe #1 relating
                                                                        Maize polymorphic Brassica sp. polym
                                                                                                        Quadruplex DNA
                                                                                                             Polynucleotide seq
                                                                                                                  Polynucleotide seq
                                                                                                                        T. brucei
                                                                                                                             Maize polymorphic
                                                                                                                                  Foldback triplex
                                                                                                                                         Foldback triplex
                                                                                                                                                  Probe Sq918 to ena
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 Thermal amplificat
                             Oligo used
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                                                         Human
                                                                                   Vector pVAC1
                                                                                        Vector
                  Oligonucleotide
                        Primer #3 for tumo
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             primer for Hum
       nuclear
                                        Machado-Jose
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                                                                                                        SEQ
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ALIGNMENTS

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RESULT 1
AAD09047/c
ID AAD090
XX AAD090
AC AAD090
XX AAD090
XX Human
XX Human;
KW Human;
KW Huntin
KW Human
KW Huma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; converse system aging: neurodegenerative disease; immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nervous system aging; neurodegenerative disease; immunological dismalignant glioma; medulloblastoma; neuroectodermal tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide HOG-403 used to construct pKS285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD09047 standard; DNA; 50 BP.
                                                                                                                                          02-NOV-2000; 2000WO-US30405.
                                                                                                                                                                                                                                                                                                                                                                                        WO200134654-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            extracellular signalling protein; HOG-403; ss.
                                                                                                                                                                                                                                                                        17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                          99US-0164025
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The present invention relates to hedgehog fusion proteins. Hedgehog contributions are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in convertebrates. Hedgehog fusion protein is useful for the prophylaxis or continuent of any condition or disease states for which a hedgehog or constituents or conditions of disease states in biological systems or conditions of disease states in biological systems. CC specimens and for diagnostic purposes in non-physiological systems. CC Hedgehog fusion protein is useful for treating neurological conditions. CC Hedgehog fusion protein is useful for treating neurological systems. CC constituents of nervous system, including Alzheimer's disease, conditions. CC parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis conditions of chronic meurodegenerative diseases of nervous system including multiple cc and chronic immunological diseases of nervous system including multiple cc sclerosis and malignant gliomas, medulloblastomas, neuroectodermal cc tumours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-403 used to construct pKS285, pKS286, pKS309, pKS310, pKS287 pKS300 and pKS289 plasmids which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 60; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001-329075/34.
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RESULT 2
AAD09078/c
                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                 Matches
Human oligonucleotide HOG-803 used to construct pMMC22.
                                                                        AAD09078 standard; DNA; 50 BP
                          04-SEP-2001
                                                                                                                                                      272 CCTTTAGCCTACAAGCAGTTTATCCCCCAATGTGGCCGAGAAGACCCTAGG 321
                                                                                                                                   50
                                                                                                                                  50;
                                                                                                                                                                                             Similarity
                                                                                                                                                                                  Conservative
                          (first entry)
                                                                                                                                                                                            100.0%;
                                                                                                                                                                                   0;
                                                                                                                                                                                            Score 50;
Pred. No.
                                                                                                                                                                                  Mismatches
                                                                                                                                                                                   0;
                                                                                                                                                                                     Indels
                                                                                                                                                                                       0;
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멍 QУ

Sequence 50 BP; 11 A; 11 C; 15 G; 13 T; 0 other;

3.2%;

DB 22;

Length .50;

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Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer; extracellular signalling protein; HOG-803; ss.
                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                              WO200134654-A1
(BIOJ ) BIOGEN INC
                                                                    05-NOV-1999;
                                                                                                                                         02-NOV-2000;
                                                                                                                                                                                                          17-MAY-2001.
                                                                                                                                         2000WO-US30405
                                                                        9905-0164025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD09079/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to hedgehog fusion proteins. Hedgehog CC proteins are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or patched protein constituent is efficacious and in the diagnosis of CC constituents or conditions of disease states in biological system or CC specimens and for diagnostic purposes in non-physiological system or CC specimens and for diagnostic purposes in non-physiological system or CC the diagnostic purposes in non-physiological systems. CC due to injury, aging of nervous system, including Alzheimer's disease, CC chronic neurodegenerative diseases of the nervous system, including Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple CC and chronic immunological diseases of nervous system including multiple cumours and to specifically target medical therapias against cancers and tumours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-803 used to construct pMMC22 plasmid which is wead in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                     05-NOV-1999;
                                                                                                                                                                                                                                                                                cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer;
                                                 (BIOJ ) BIOGEN INC.
                                                                                                                       02-NOV-2000; 2000WO-US30405.
                                                                                                                                                               17-MAY-2001
                                                                                                                                                                                                  WO200134654-A1.
                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                        Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injur
                                                                                                                                                                                                                                                                                                                                                                                               Human oligonucleotide HOG-804 used to construct pMMC23.
                                                                                                                                                                                                                                                                    extracellular signalling protein; HOG-804; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                         04-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 50 BP; 6 A; 17 C; 13 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD09079 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 62; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         multiple sclerosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-329075/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Strauch K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                 99US-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%; 98.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD09081/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to hedgehog fusion proteins. Hedgehog CC proteins are a family of extracellular signalling proteins that regulate CC various aspects of embryonic development both in vertebrates and in CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or CC patched protein constituent is efficacious and in the diagnosis of CC constituents or conditions of disease states in biological system or CC specimens and for diagnostic purposes in non-physiological systems. CC Hedgehog fusion protein is useful for treating neurological systems. CC Hedgehog fusion protein is useful for treating neurological conditions can be considered to injury, aging of nervous system, including Alzheimer's disease, CC chronic neurodegenerative diseases of the nervous system, including CC and chronic immunological diseases of the nervous system including multiple CC and chronic immunological diseases of nervous system including multiple colorosis and malignant gliomas, medulloblastomas, neuroectodermal CC tumours and to specifically target medical therapies against cancers and CC is human oligonucleotide HOG-804 used to construct pMMC23 plasmid which cc is used in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                            cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; nervous system aging; neurodegenerative disease; immunological disease;
(BIOJ ) BIOGEN INC
                                       05-NOV-1999;
                                                                          02-NOV-2000; 2000WO-US30405
                                                                                                                                                                                                                                                          malignant glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                        17-MAY-2001
                                                                                                                                                                WO200134654-A1.
                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                         extracellular signalling protein; HOG-807; ss.
                                                                                                                                                                                                                                                                                                                                       Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant;
                                                                                                                                                                                                                                                                                                                                                                                   Human oligonucleotide HOG-807 used to construct pMMC26.
                                                                                                                                                                                                                                                                                                                                                                                                                                   04-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD09081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD09081 standard; DNA; 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 50 BP; 6 A; 19 C; 13 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             228 CGGGCAGGGGTTCGGGAAGAGGAGGCACCCCAAAAAGCTGACCCCTTTA 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    is used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 62; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-329075/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50 CGGGCAGGGGGTTCGGGAGGAGGAGGCACCCCAAAAAGCTGACCCCTTTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                         99US-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 48.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ببر
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Novel isolated hedgehog fusion polypeptide useful for treating

WPI; 2001-329075/34.

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RESULT 5
AAD09073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cc various aspects of embryonic development both in vertebrates and in cc invertebrates. Hedgehog fusion protein is useful for the prophylaxis or cc treatment of any condition or disease state for which a hedgehog or cc patched protein constituent is efficacious and in the diagnosis of constituents or conditions of disease states in biological system or cc specimens and for diagnostic purposes in non-physiological system or cc specimens and for diagnostic purposes in non-physiological systems. Cc Hedgehog fusion protein is useful for treating neurological conditions cc chronic neurodegenerative diseases, including Alzheimer's disease, cc chronic neurodegenerative diseases of the nervous system, including Cc parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple csclerosis and malignant gliomas, meduloblastomas, neuroectodermal cc sclerosis and malignant gliomas, meduloblastomas, neuroectodermal cc timours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-807 used to construct pMMC26 plasmid which
밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neurological conditions
disease, Huntington's cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that regulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 62; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                              Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer; extracellular signalling protein; HOG-789; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 50 BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide HOG-789 used to construct pMMC22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD09073 standard; DNA; 47
            Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                  WPI; 2001-329075/34.
                                                                                                                                                                           (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                      05-NOV-1999;
                                                                                                                                                                                                                                                          02-NOV-2000;
                                                                                                                                                                                                                                                                                                  17-MAY-2001
                                                                                                                                                                                                                                                                                                                                         WO200134654-A1
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                228 CGGGCAGGGGTTCGGGAAGAGGGAGGCACCCCAAAAAGCTGACCCCTTTA 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50 CGGGCAGGGGGTTCGGGAAGAGGAGCACCCCCAGCAGCTGACCCCTTTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 Similarity 47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                              2000WO-US30405.
                                                                                                                                                                                                                      9908-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 C; 15 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ns such as Alzheimer's disease, Parkinson's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 45.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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The present invention relates to hedgehog fusion proteins. Hedgehog CC proteins are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or CC patched protein constituent is efficacious and in the diagnosis of CC constituents or conditions of disease states in biological systems or CC specimens and for diagnostic purposes in non-physiological systems. CC Hedgehog fusion protein is useful for treating neurological conditions CC thouse the injury, aging of nervous system, including Alzheimer's disease, CC chronic neurodegenerative diseases of the nervous system including parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple selerosis and malignant gliomas, medulloblastomas, neuroectodermal CC sclerosis and malignant gliomas, medulloblastomas, neuroectodermal tumours and to specifically target medical therapies against cancers and the summan oligonucleotide HOG-789 used to construct pMMC22 plasmid which can be immediant the incention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 61; 178pp; English.
                                                                 in the invention.
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RESULT 6
AAD09074
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                       Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer;
                                                                                                                                                                                                                             extracellular signalling protein; HOG-791;
                                                                                                                                                                                                                                                                                                                                                         AAD09074 standard; DNA; 47
                                                                                                                                                                                                                                                                                                 Human oligonucleotide HOG-791 used to construct pMMC23
                                                                                                                                                                                                                                                                                                                     04-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                        AAD09074;
                                                                                                                                                                                                                                                                                                                                                                                                                          225 GACCGGGCAGGGGGTTCGGGAAGAGGGGAGGCACCCCAAAAAGCTGACC 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                       45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 43.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson disease, Huntington's chorea, amylotropic lateral sclerosis, and the second school of the sec

Example 1; Page 61-62; 178pp; English

multiple sclerosis

WPI; 2001-329075/34.

(BIOJ) BIOGEN INC 05-NOV-1999; 02-NOV-2000; 2000WO-US30405

99US-0164025

17-MAY-2001 WO200134654-A1 Homo sapiens.

g

Sequence 47 BP; 13 A; 13 C; 18 G; 3 T; 0 other;

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RESULT 7
AAD09077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                            Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's
                                                                                                    multiple sclerosis
                                                                                                                                                                                       WPI; 2001-329075/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer; extracellular signalling protein; HOG-799; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis;
                                                                                                                                                                                                                                                        (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                          02-NOV-2000; 2000WO-US30405.
                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human oligonucleotide HOG-799 used to construct pMMC22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment of any condition or disease state for which a hedgehog or patched protein constituent is efficacious and in the diagnosis of constituents or conditions of disease states in biological system or specimens and for diagnostic purposes in non-physiological systems. Hedgehog fusion protein is useful for treating neurological conditions due to injury, aging of nervous system, including Alzhelmer's disease, chronic neurodegenerative diseases of the nervous system, including Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple sclerosis and malignant gliomas, medulloblastomas, neuroectodermal tumours and to specifically target medical therapies against cancers and the suppose the receptor for the protein. The present sequence is human oligonucleotide HOG-791 used to construct pMMC23 plasmid which is used in the invention.
                                                                                                                                                                                                                                                                                                                                                               17-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD09077 standard; DNA; 45 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 47 BP; 11 A; 13 C; 20 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in invertebrates. Hedgehog fusion protein is useful for the prophylaxis or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                        Huntington's chorea, amylotropic lateral sclerosis, and sclerosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                        9905-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 43.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . No. 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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various aspects of embryonic development both in vertebrates and in The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that reg

regulate

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene

62; 178pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF27031/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or CC treatment of any condition or disease state for which a hedgehog or CC patched protein constituent is efficacious and in the diagnosis of CC constituents or conditions of disease states in biological system or CC specimens and for diagnostic purposes in non-physiological systems. CC Hedgehog fusion protein is useful for treating neurological conditions CC due to injury, aging of nervous system, including Alzheimer's disease, CC chronic neurodegenerative diseases of the nervous system, including CC arkinson's disease, Huntington's chorea, amylotropic lateral sclerosis CC and chronic immunological diseases of nervous system including multiple CC sclerosis and malignant gliomas, medulloblastomas, neuroectodermal CC tumours and to specifically target medical therapies against cancers and CC tumours which express the receptor for the protein. The present sequence is human oligonucleotide Hog-799 used to construct pMMC22, pMMC23, pMMC25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                 Example 6; Page 77; 157pp; English.
                                                                                                 polyalkylene glycol group linked to any residue other than N-terminal and lysine residues - \,
                                                                                                                               Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a
                                                                                                                                                                                           WPI; 2001-049927/06.
                                                                                                                                                                                                                                                                                                                     01-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder;
                                                                                                                                                                                                                                                               (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                          07-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                malignant glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF27031 standard; DNA; 48
                                                                                                                                                                                                                                                                                                    13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                       26-MAY-2000; 2000WO-US14741.
                                                                                                                                                                                                                                                                                                                                                                                                                            WO200073337-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mutagenic primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammatory disorder; autoimmune disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 45 BP; 12 A; 14 C; 10 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and pMMC26 plasmids which are used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      272 CCTTTAGCCTACAAGCAGTTTATCCCCAATGTGGCCGAGAAGACC 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CCTTTAGCCTACAAGCAGTTTATCCCCAAGGTGGCCGAGAAGACC 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                           Taylor F,
                                                                                                                                                                                                                                                                                                  99US-0137011
99US-0149016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%;
97.8%;
                                                                                                                                                                                                                           Garber E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 43.4;
Pred. No. 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0;

glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of

hedgehog

WPI; 2001-049927/06.

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defining and mapping functionally important regions of a protein by comodifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of comparison of a protein of the conformation of the protein serious medical conditions including various neurological disorders. Comparison of a protein of comparison of a protein of the conformation of the management of comparison of the protein of the management of comparison of the inversing or ameliorate neurological disease, and cancers. In particular, they compare used to prevent preventing or ameliorate neurological injury and concerns of trauma; immunological diseases, altaheimer's conformation of the inverse of the nervous system (e.g., multiple conformation); stroke; and malignant gliomas, medulloblastomas and conformation in increased shallity in solutions made to the hedgehog protein conformation to cross the blood-brain from protecolytic conformation of the invention to cross the blood-brain barrier, and an conformation of the invention to protein the present sequence represents a human Sonic hedgehog mutagenic primer used in an exemplification of the invention of the primer used in an exemplification of the invention of the primer used in an exemplification of the conformal stability multiple and an advantage when formulating the human Sonic hedgehog mutagenic primer used in an exemplification of the conformal stability multiple and an exemplification of the conformal stability multiple and an exemplification of the conformal stability and the primer used in an exemplification of the conformal stability and the primer used in an exemplification of the conformal stability and the primer used in an exemplification of the conformal stability and the primer used in an exemplification of the conformal stability and the primer used in an exemplification of the conformation to cross the blood-primer and an exemplification of 
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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 48 BP; 10 A; 13 C; 15 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF27034;
                Pepinsky RB,
                                                                (BIOJ ) BIOGEN INC
                                                                                                            01-JUN-1999;
13-AUG-1999;
                                                                                                                                                                               26-MAY-2000; 2000WO-US14741.
                                                                                                                                                                                                                          07-DEC-2000
                                                                                                                                                                                                                                                                           WO200073337-A1
                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                             malignant glioma;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          neurodegenerative disorder; Parkinson's disease; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF27034 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    278 GCCTACAAGCAGTTTATCCCCAATGTGGCCGAGAAGACCCTAGGCGCC 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                   Taylor F,
                                                                                                              99US-0149016
                                                                                                                                      99US-0137011
                                                                                                                                                                                                                                                                                                                                                                                                               medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%;
93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 43.2;
Pred. No. le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 10.0); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
. 1e+02;
. 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22; Length 48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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cc which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene cc glycol group, with the proviso that the polymer is not conjugated to the conjugated not the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog consisting an edgehog (Ihh), or Desert hedgehog (protein used in the conjugate may be a wild-type or mutant Sonic hedgehog consisting and mapping functionally important regions of a protein by commodifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoinmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative conjugates (e.g., Parkinson's disease, Huntington's disease, Alzheimer's cisease); age-associated neurological disease; neurological injury and crauma; immunological diseases of the nervous system (e.g., multiple selerosis); stroke; and malijanat gliomas, medulloblastomas and cancercidermal tumours. The modifications made to the hedgehog protein of the vasculature for prolonged periods may acid on from proteolytic degradation, or reduced immunogenicity. In particular, the ability to stability in solution, protection from proteolytic degradation, the vasculature for prolonged periods may allow a hedgehog of protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the chedgehog protein of the degehod mutagenic provised to an advantage when formulation of the hedgehod mutagenic provised to an average when formulation of the medgehod mutagenic provised to an accentification of the medgehod mutagenic provised to an accentification of the medgehod mutagenic provised to a second provis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-terminal and lysine residues
                                                                                         Sonic hedgehog mutagenic primer used in an exemplification
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Вþ QУ Matches Query Match 398 GACATCATATTTAAGGATGAAGAAAACACCGGAGCGGACAGGCTGATG 445 Local Similarity Conservative 2.7%; 93.8%; 0; Mismatches Score 43.2; DB 22; Pred. No. le+02; ω •• Length 48; Indels 0; Gaps 0

Sequence 48 BP; 10 A;

15 C; 7 G; 16 T; 0 other;

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AAD09080/c
                                                                                                                                                                                                                                                                 RESULT 10
Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant;
                                                    Human oligonucleotide HOG-806 used to construct pMMC25.
                                                                                                         04-SEP-2001
                                                                                                                                                                                                                  AAD09080 standard; DNA; 50
                                                                                                           (first entry)
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cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; extracellular malignant glioma; medulloblastoma; nervous system aging; neurodegenerative disease; immunological disease; neuroectodermal tumour;

PDD XXX WO200134654-A1

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RESULT 11
AAD09082/c
ID AAD090
XX AAD090
XX AAD090
XX Human
DT 04-SEP
XX Human;
KW Human;
KW Huntin
KW nervou
KW mervou
KW malign
KW malign
XX Homo s
XX WO2001
XX WO2001
XX WO2001
XX PP 17-MAY
XX PF 02-NOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to hedgehog fusion proteins. Hedgehog CC proteins are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in CC invertebrates. Hedgehog fusion protein is useful for the prophylaxis or CC patched protein constituent is efficacious and in the diagnosis of CC constituents or conditions of disease states in biological system or CC specimens and for diagnostic purposes in non-physiological systems. CC Hedgehog fusion protein is useful for treating neurological systems. CC thronic neurodegenerative diseases of the nervous system, including Alzheimer's disease, CC chronic neurodegenerative diseases of the nervous system including CC and chronic immunological diseases of the nervous system including CC and chronic immunological diseases of nervous system including multiple collections and to specifically target medical therapies against cancers and tumours and to specifically target medical therapies against cancers and construct which express the receptor for the protein. The present sequence collection in the invention
                                                                                                                                                                      nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer; extracellular signalling protein; HOG-808; ss.
                                                                                                                                                                                                                            Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis;
              02-NOV-2000; 2000WO-US30405
                                                         17-MAY-2001
                                                                                                    WO200134654-A1
                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                   Human oligonucleotide HOG-808 used to construct pMMC22.
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD09082 standard; DNA; 43 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 50 BP; 6 A; 16 C; 16 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 62; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                228 CGGGCAGGGGTTCGGGAAGAGGGAGGCACCCCAAAAAGCTGACCCCTTTA 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-NOV-2000; 2000WO-US30405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50 CGGGCAGGGGGTTCGGGCAGCAGCAGCACCCCAAAAAGCTGACCCCTTTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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AAD09076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that regulate Various aspects of embryonic development both in vertebrates and in invertebrates. Hedgehog fusion protein is useful for the prophylaxis or treatment of any condition or disease state for which a hedgehog or patched protein constituent is efficacious and in the diagnosis of constituents or conditions of disease states in biological system or specimens and for diagnostic purposes in non-physiological systems. Hedgehog fusion protein is useful for treating neurological conditions due to injury, aging of nervous system, including Alzheimer's disease, chronic neurodegenerative diseases of the nervous system, including multiple sclerosis and chronic immunological diseases of nervous system including multiple sclerosis and malignant gliomas, medulloblastomas, nervoectodermal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                   05-NOV-1999;
                                                                                                                                                                                                                         nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer; extracellular signalling protein; HOG-797; ss.
                                                                                                                                                                                                                                                                                 Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant; cytostatic; therapy; Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis;
                                                                    02-NOV-2000; 2000WO-US30405
                                                                                                                17-MAY-2001
                                                                                                                                                      WO200134654-A1.
                                                                                                                                                                                                                                                                                                                                                                 Human oligonucleotide HOG-797 used to construct pMMC26.
                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                             04-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD09076 standard; DNA; 47 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 43 BP; 8 A; 10 C; 14 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and pMMC26 plasmids which are used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumours and to specifically target medical therapies against cancers and tumours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-808 used to construct pMMC22, pMMC23, pMMC25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 62; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-329075/34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              278 GCCTACAAGCAGTTTATCCCCAATGTGGCCGAGAAGACCCTAG 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43 GCCTACAAGCAGTTTCTCCCCAATGTGGCCGAGAAGACCCCTAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                     99US-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0164025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.6%;
97.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 41.4;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .1e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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(BIOJ) BIOGEN INC

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invertebrates. Hedgehog fusion protein is useful for the prophylaxis or treatment of any condition or disease state for which a hedgehog or patched protein constituent is efficacious and in the diagnosis of constituents or conditions of disease states in biological system or specimens and for diagnostic purposes in non-physiological systems. Hedgehog fusion protein is useful for treating neurological conditions due to injury, aging of nervous system, including Alzheimer's disease, chronic neurodegenerative diseases of the nervous system, including parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sclerosis and malignant gliomas, medulloblastomas, neuroectodermal tumours and to specifically target medical therapies against cancers and tumours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-797 used to construct pMMC23 plasmid which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that regulate various aspects of embryonic development both in vertebrates and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated hedgehog fusion polypeptide useful for treating neurological conditions such as Alzheimer's disease, Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strauch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 62; 178pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-329075/34
                                                                                                                                                                                                                                                                                       cytostatic; therapy, Alzheimer's disease; Parkinson's disease; injury; Huntington's chorea; amylotropic lateral sclerosis; multiple sclerosis; nervous system aging; neurodegenerative disease; immunological disease; malignant glioma; medulloblastoma; neuroectodermal tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 47 BP; 9 A; 15 C; 20 G; 3 T; 0 other;
                                                    (BIOJ ) BIOGEN INC
                                                                                        05-NOV-1999;
                                                                                                                           02-NOV-2000; 2000WO-US30405.
                                                                                                                                                                17-MAY-2001
                                                                                                                                                                                                    WO200134654-A1
                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                      extracellular signalling protein; HOG-795; ss
                                                                                                                                                                                                                                                                                                                                                                 Human; hedgehog protein; nootropic; neuroprotective; anticonvulsant;
                                                                                                                                                                                                                                                                                                                                                                                                       Human oligonucleotide HOG-795 used to construct pMMC25 plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                            04-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD09075;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD09075 standard; DNA; 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             225 GACCGGGCAGGGGGTTCGGGAAGAGGGAGGCACCCCAAAAAGCTGACC 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ш
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                        99US-0164025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 40.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    patched protein constituent is efficacious and in the diagnosis of constituents or conditions of disease states in biological system or specimens and for diagnostic purposes in non-physiological systems. Hedgehog fusion protein is useful for treating neurological conditions disease, against the injury, aging of nervous system, including Alzheimer's disease, chronic neurodegenerative diseases of the nervous system, including Parkinson's disease, Huntington's chorea, amylotropic lateral sclerosis and chronic immunological diseases of nervous system including multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to hedgehog fusion proteins. Hedgehog proteins are a family of extracellular signalling proteins that regulativarious aspects of embryonic development both in vertebrates and in invertebrates. Hedgehog fusion protein is useful for the prophylaxis or treatment of any condition or disease state for which a hedgehog or treatment of any condition or disease state for which a hedgehog or treatment of any condition or disease state for which a hedgehog or treatment of any condition or disease state for which a hedgehog or treatment of any condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition or disease state for which a hedgehog or treatment of the condition of th
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sclerosis and malignant gliomas, medulloblastomas, neuroectodermal tumours and to specifically target medical therapies against cancers and tumours which express the receptor for the protein. The present sequence is human oligonucleotide HOG-795 used to construct pMMC25 plasmid which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 62; 178pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  multiple scierosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-329075/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 47 BP; 11 A; 16 C; 17 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF27032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF27032 standard; DNA; 42 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is used in the invention.
                                                                                              (BIOJ ) BIOGEN INC.
                                                                                                                                                                                                01-JUN-1999;
                                                                                                                                                                                                                                                                26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                          WO200073337-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mutagenic primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   malignant glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                225 GACCGGGCAGGGGGTTCGGGAAGAGGGAGGCACCCCAAAAAGCTGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                2000WO-US14741.
                                                                                                                                                                     99US-0149016
                                                                                                                                                                                                99US-0137011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.4%;
87.2%;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 37.4; DB 22; Pred. No. 9.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47;
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Pepinsky RB,

Taylor F,

Garber

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RESULT 15
AAF27036/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc a hedgehog fusion protein. The invention also relates to methods of combifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the position and/or type of modification have on the activity of the comprosed methods in the protein method of the comprosed methods in the management of comprosed methods in the methods in the management of comprosed methods in the methods of comprosed methods in the methods may allow a hedgehog comprosed methods in the methods may allow a hedgehog comprosed methods in the methods may allow a hedgehog comprosed methods in the methods may allow a hedgehog comprosed methods in the methods may allow a hedgehog comprosed methods in the methods may allow a hedgehog comprosed methods in the methods of comprosed methods in the methods in the methods in the methods may allow a medical methods in the 
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                                                                                                                                                                              bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease;
                                                Synthetic
                                                                        Homo sapiens
                                                                                                                     malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss.
                                                                                                                                                        Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                           Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                                                                                                                                                                                                                                                                              Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:40.
                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF27036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF27036 standard; DNA; 42 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 42 BP; 8 A; 13 C; 9 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                338 GAAGGGAAGATCTCCAGAAACTCCGAGCGATTTAAGGAACTC 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-terminal and lysine residues -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-049927/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.4%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 37.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ω,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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XXX
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AAF27039/c
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                                                                                                                                                                                                                                                                                                               Qy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              various medical conditions including various neurological disorders,

C inflammatory and autoimmune diseases, and cancers. In particular, they

may be used to prevent preventing or ameliorate neurodegenerative

C disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's

C disease); age-associated neurological disease; neurological injury and

C trauma; Immunological diseases of the nervous system (e.g., multiple

C sclerosis); stroke; and malignant gliomas, medulloblastomas and

C neuroectodermal tumours. The modifications made to the hadgehog protein

C may result in increased half-life, altered tissue distribution (such as

C may result ability to stay in the vasculature for longer periods of

C degradation, or reduced immunogenicity. In particular, the ability to

C degradation, or reduced immunogenicity. In particular, the ability to

C momain in the vasculature for prolonged periods may allow a hedgehog

C protein of the invention to cross the blood-brain barrier, and an

C increased thermal stability would be an advantage when formulating the

C human sonic hedgehog multagenic primer used in an exemplifications of the
                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
  Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:43.
                                                        30-MAR-2001 (first entry)
                                                                                                                                                AAF27039 standard; DNA; 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog protein. The hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of a hedgehog functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various madical accordance in changing various may be used in the management of
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 42 BP; 11 A; 14 C; 9.G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                   474 CTTTGGCCATCTCGGTGATGAACCAGTGGCCAGGAGTGAAAC 515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human Sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-terminal and lysine residues -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-049927/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pepinsky RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOJ ) BIOGEN INC.
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                                                                                                                                                                                                                                                           42 CTTTGGCCATCTCGGTGATGTGTCAGTGGCCAGGAGTGAAAC
                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                         39;
                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Taylor F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0149016
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                                                                                                                                                                                                                                                                                                                                                                          2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Garber E;
                                                                                                                                                                                                                                                                                                                                                                          Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                          Score 37.2;
                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                          1e+03;
                                                                                                                                                                                                                                                                                                                                                                                             DB 22;
                                                                                                                                                                                                                                                                                                                                                  ω
                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                          Length 42;
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                                                                                                                                                                                                                                                                                                                                             Gaps
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QΥ
                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease); age-associated neurological disease; neurological injury and disease; immunological diseases of the nervous system (e.g., multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Modified hedgehog protein, useful in the treatment of Parkinson' disease and Huntington's chorea, comprises a polymer containing polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammatory disorder; autoimmune disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the provison that the polymer is not conjugated to the
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                                                                                                                                                                                                                                                                                                                                                                                                     sclerosis); stroke; and malignant gliomas, meduiloblastomas and neuroectodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog manufacture for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated in the vasculature for prolonged periods may allow a hedgehog treated may be a provinced in the vasculature for prolonged periods may be a provinced in the vasculature for prolonged periods may be a provinced in the vasculature for prolonged periods may be a provinced in the vasculature for prolonged periods may be a provinced in the vasculature for prolonged periods may be a provinced in the vasculature for prolonged periods may allow a hedgehog periods may allow a hedgehog periods may be a provinced in the vascu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 6; Page 77; 157pp; English.
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                                                                                                                                                                                            Sequence 38
                                                                                                                                                                                                                                                                                 increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human Sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                    protein of the invention to cross the blood-brain barrier, and an
      662 GACTGGGTGTACTACGAGTCCAAGGCACATATCCACTG
                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001-049927/06
                                                                                                      Similarity
                                                                                                                                                                                                  BP; 8
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Taylor
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                                                                                                                                                                                                  A; 11 C;
                                                                                                   2.3%;
                                                                                                                                                                                                      9 G;
                                                                           0;
                                                                                                      Pred.
                                                                                                            Score 36.4;
Pred. No. 1
                                                                                                                                                                                                      10 T; 0 other;
                                                                               Mismatches
                                                                                                            .4e+03;
                                                                                                                                             DB
                                                                                                                                             Length
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                                                                                  Gaps
                                                                                     0;
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В

38

GACTGGGTGTACTACGAGTGCAAGGCACATATCCACTG

hedgehog protein

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RESULT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mutagenic primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF27025;
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                                                         inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative adisorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease); age-associated neurological disease; neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, medulloblastomas and protein neuroectodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog remain in the vasculature for prolonged periods may allow a hedgehog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        malignant glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bioavailibility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sonic hedgehog; Shh; polymer conjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAR-2001
                                                                                                                                                                                                                                                                                                                         a neugrino tustom protein by defining and mapping functionally important regions of a protein by defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they
                                                                                                                                                                                                                                                                                                                                                                                                                                                     conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the glycol group, with the proviso that the polymer is not conjugated to the y-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease and Huntington's chorea, comprises a polymer containing polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-1999;
13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-2000; 2000WO-US14741.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200073337-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to novel polymer conjugates of hedgehowhich have increased bioavailability. The hedgehog proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-terminal and lysine residues -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified hedgehog protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-049927/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOJ ) BIOGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pepinsky RB,
                                           protein of the invention to cross the blood-brain barrier, and
                          ncreased thermal stability would be an advantage when formulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 67; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Taylor F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0137011
99US-0149016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            onn; polymer conjugate; polyalkene glycol group; formulation; neurological disorder;
  in powder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Garber E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful in the treatment of Parkinson's
       form. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprises a polymer containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             conjugates of hedgehog proteins
       sequence represents
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RESULT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Phh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of inflammatory and autolimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's
                                                                                                                                                                                                                                The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are
                                                                                                                                                                                                                                                                                                              Modified hedgehog protein, useful in the treatment of Parkinson' disease and Huntington's chorea, comprises a polymer containing polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues -
                                                                                                                                                                                                                                                                                   Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            maiignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Pepinsky RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-1999;
13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sonic hedgehog; Shh; polymer conjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOJ ) BIOGEN INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bioavailibility;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF27038 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human Sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TUACTOU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 703 GGTGAAAGCAGAGAACTCGGTGGCGGCCAAATCGGGAGGCTGCT 746
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Taylor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0137011
99US-0149016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39
                                                                                                                                                                                                                                                                                                                                                                                                                                    Garber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 36;
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Best Local :
                     The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are
                                                                                                                                              Modified hedgehog protein, usefu disease and Huntington's chorea,
                                                                                 Example 6;
                                                                                                                                                                                                                                                                                                         01-JUN-1999;
13-AUG-1999;
                                                                                                               N-terminal and lysine residues
                                                                                                                               polyalkylene glycoľ group linked
                                                                                                                                                                                                      WPI; 2001-049927/06
                                                                                                                                                                                                                                      Pepinsky RB,
                                                                                                                                                                                                                                                                      (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease); age-associated neurological disease; neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, medulloblastomas and neuroectodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the human sonic hedgehog mutagenic primer used in an exemplification of the invention of the stability would be an advantage.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mutagenic primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neurodegenerative disorder; Parkinson's disease; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             malignant glioma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF27035;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        597 CCACGTCTGACCGCGACCGCAGCAAGTACGGCATGCTGG 635
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCACGTCTGACCGCGATCGCTGCAAGTACGGCATGCTGG 1
                                                                               Page 77; 157pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                      Taylor F,
                                                                                                                                                                                                                                                                                                         99US-0149016
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         medulloblastoma; neuroectodermal tumour;
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                                                                                                                                        useful in the treatment of Parkinson's horea, comprises a polymer containing a
                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 35.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                               ţ
                                                                                                                               any residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0,

conjugated to a non-naturally-occurring polymer comprising a polyalky glycol group, with the proviso that the polymer is not conjugated to

comprising a polyalkylene

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DRXX REPRESENTATION OF STREET CONTRACTOR OF STREET 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease); age-associated neurological disease; neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, medulloblastomas and neuroectodermal tumours. The modifications made to the hedgehog protein result in increased half-life, altered tissue distribution (such as may result in increased half-life, altered tissue distribution (such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          various medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 42 BP; 11 A; 11 C; 10 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF27037;
WPI; 2001-049927/06
                                                                 Pepinsky RB,
                                                                                                                                                                                                                    01-JUN-1999;
13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             07-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200073337-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               malignant glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurodegenerative disorder; Parkinson's disease; Huntington's disease;
Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF27037 standard; DNA; 37 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human Sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nedgehog protein in powder form. The present sequence represents
                                                                                                                                              (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                26-MAY-2000; 2000WO-US14741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 445 GACTCAGAGGTGTAAGGACAAGTTGAACGCTTTGGCCCATCTC 486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GACTCAGAGGTGTAAAGGACTGCTTAAACGCTTTGGCCATCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                     Taylor
                                                                                                                                                                                                                    99US-0149016
                                                                                                                                                                                                                                                          99US-0137011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 35.6;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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conjugated to a non-naturally-occurring polymer compusates of hedgehog proteins are glycol group, with the proviso that the polymer is not conjugated to the conjugated to the proviso that the polymer is not conjugated to the conjugate may be a wild-type or mutant Sonic hedgehog (Protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dh) protein. The hedgehog conditions and mapping functionally important regions of a protein by composition and/or type of modification have on the activity of the modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog protein by composition and/or type of modification have on the activity of the protein. The hedgehog protein by composition and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of compositions medical conditions including various neurological disorders, inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurological injury and creased (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease; age-associated neurological disease; neurological injury and creased the protein diseases of the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, meduliblastomas and result in increased half-life, altered trissue distribution (such as an improved ability to stay in the vasculature for longer periods of the hedgehog protein of the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an conjugate of the hedgehog multagenic primer used in an exemplification of the human sonic hedgehog multagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's Chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 77; 157pp; English
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g
                                              Matches
                                                                   Query Match
                      538 CGAAGATGGCCACCACTCAGAGGAGTCTCTGCACTAC 574
37 CGAAGATGGCCACCACTGCGAGGAGTCTCTGCACTAC
                                                          Local
                                                           Similarity
                                              Conservative
                                                          2.18; 94.68;
                                                 Mismatches
                                                             Score 33.8;
Pred. No. 3
                                                           Pred.
                                                             .9e+03
                                                 Indels
                                                                       Length 37;
                                                  0;
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Sequence 37

BP; 6 A; 10 C; 12 G; 9 T; 0 other;

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RESULT 21
AAF27041/
Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:45
                                                                                     30-MAR-2001
                                                                                                AAF27041;
                                                                                                          AAF27041 standard; DNA; 35 BP
                                                                                     (first entry)
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inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
Homo
                                                                          mutagenic
                                                                                                                                                                                                                                                                       bioavailibility; formulation; neurological disorder;
                                                                                                                                                                                                                                                                                                             Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                                                                                   medulloblastoma; neuroectodermal tumour;
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WO200073337-A1

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AAF27040/c
ID AAF270
XX
AC AAF270
XX
DT 30-MAR
XX
DE Human
XX
KW Sonic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ωy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;
                                                              Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease); age-associated neurological disease; neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple neuroectodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for proteolytic remain in the vasculature for proteols and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a term of the invention of the present sequence represents a series of the book of the invention of the present sequence represents a series of the book of the present sequence represents a series of the book of the present sequence represents a series of the book of the present sequence represents a series of the present sequence represents and the present sequence represents a series of the present sequence and the present sequence and
                                                                                                                                         30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a hedgehog fusion protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of inflammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (or Darviness) and cancers. In particular, they have described the conditions including or ameliorate neurodegenerative.
                                                                                                                                                                                                                                                         AAF27040 standard; DNA; 37 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 35 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human Sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog
                                                                                                                                                                                                                                                                                                                                                                                                                                                           639 GCCTGGCGGTGGAGGCCGGCTTCGACTGGGTGTAC 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-049927/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-terminal and lysine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-MAY-2000; 2000WO-US14741.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                      35 GCCTGGCGGTGGAGGCCTGCTTCGACTGGGTGTAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be
                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Taylor F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0149016.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0137011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 15 C; 9 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.1%; 97.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Garber E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 33.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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믕 Qy

538 CGAAGATGGCCACCACTCAGAGGAGTCTCTGCACTAC 574

37 CGAAGATGGCCACCACTCATGCGAGTCTCTGCACTAC 1

Matches

Local

Similarity

2.0%;

Conservative

0; Pred. Score 32.2; ed. No. 7.2 Mismatches

7.2e+03 DB 22;

Length 37; Indels

0;

Gaps

0;

Sequence 37 BP; 7 A;

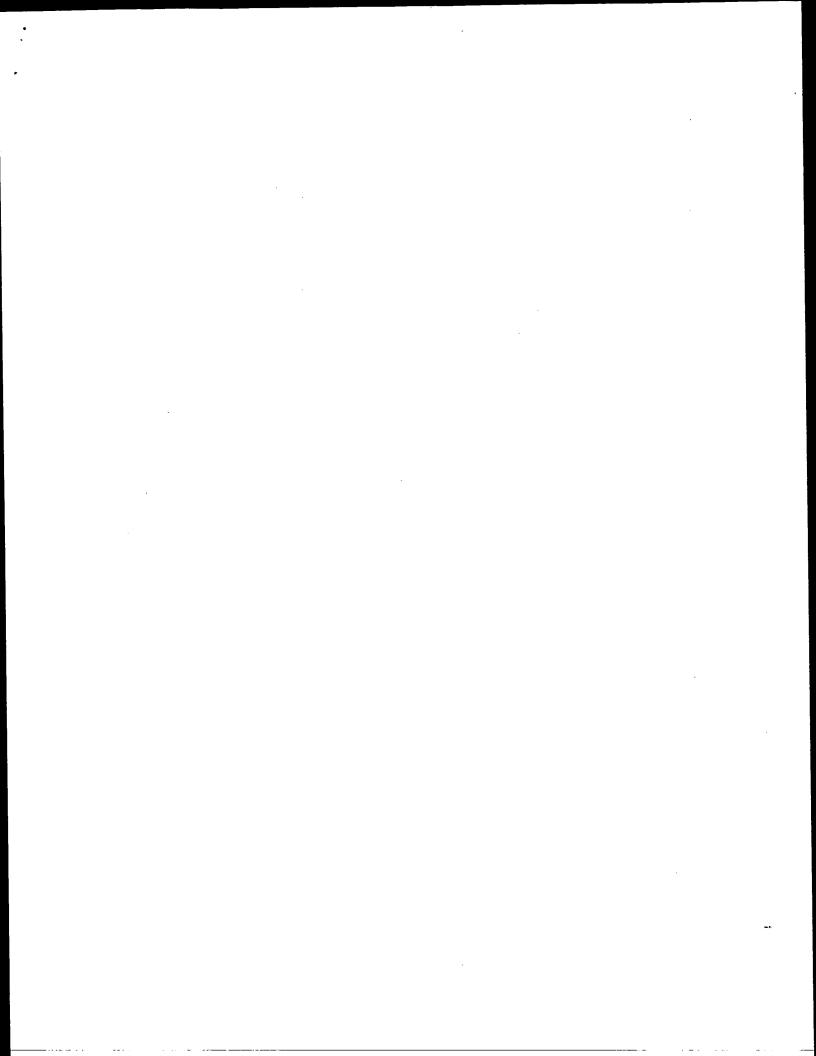
8 C; 13 G; 9 T; 0 other;

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defining and mapping functionally important regions of a protein by comodifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the comodifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the comodification including various neurological disorders (comparison medical conditions including various neurological disorders, comparison and cancers. In particular, they may be used to prevent preventing or ameliorate neurological injury and disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's comparison of the neurological disease, and cancers in the proventing or ameliorate neurological injury and crauma; immunological diseases, Huntington's disease, Alzheimer's comparison of the neurological disease, and cancers in the proventing or ameliorate neurological injury and comparison in the vasculation and malignant gliomas, meduloblastomas and comparison in the vasculations and the hedgehog protein of the invention to cross the board-may allow a hedgehog comparison of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the comparison of the invention in the present sequence represents a comparison of the invention of the protein of the protein of the invention of the protein sequence represents a comparison of the invention of primer used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               YX PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PDX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 77; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-terminal and lysine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-049927/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BIOJ ) BIOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAY-2000; 2000WO-US14741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200073337-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mutagenic primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bioavailibility; formulation; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         glioma; medulloblastoma; neuroectodermal tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Taylor F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0149016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0137011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Garber E;
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ABT03768/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 23
                                                                                                                                                                                   RESULT 24
AAQ33571
B
                                                                                                                                                                                                                                                               QΥ
                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for determining the presence of neoplastic molecular markers in a host, involving the use of neoplastic molecular markers pecific reagents to detect such markers and analysing the array of reagents, allowing the identification of the neoplastic disease present. This can be used to determine the best treatment for cancers, in particular neural cell, lung and prostate tumours. The present sequence is a PCR primer useful for detecting the coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human SHH gene PCR primer SEQ ID NO: 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABT03768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABT03768 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the reagents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-537346/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Palm K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-NOV-2000; 2000US-249508P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-2001; 2001WO-US43461.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transcription factor; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; cancer; neoplastic disease; tumour specific marker; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CEMI-) CEMINES LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 19; 41pp; English.
WO9213102-A
                         Bos taurus
                                                  genetic mapping;
                                                                                          Microsatellite
                                                                                                                     02-FEB-1993
                                                                                                                                              AAQ33571;
                                                                                                                                                                        AAQ33571 standard; DNA; 46
                                                                                                                                                                                                                                                                                                                                              Sequence 27 BP; 3 A; 11 C; 9 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                         sequences of markers of the invention.
                                                                                                                                                                                                                                                                  755 TCGGCCACGGTGCACCTGGAGCAGGGC 781
                                                                                                                                                                                                                                         27 TCGGCCACGGTGCACCTGGAGCAGGGC 1
                                                              selection; primers;
                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                   (first entry)
                                                                                        sequence from clone AGLA259.
                                                  traits; amplification; ss
                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                      1.7%;
                                                                OPTIPRIM; breeding;
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                                                                                                                                                                         ВP
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                                                                                                                                                                                                                                                                                                                      Score 27;
                                                                                                                                                                                                                                                                                                           Pred. No.
                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                           DB 24;
5.4e+04;
                                                                  cattle; parentage;
                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                      Length 27;
                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                0;
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AC AAF
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AC AAF
XX DA
XX DA
XX Syn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence is that of a bovine microsatellite sequence obtd. by cc screening a library of bovine MboI DNA fragments of between cc 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. CC distribution of microsatellites and MboI sites, the frequency of cc (T6)n >9 microsatellites in the bovine genome is estimated at >100, cc (T6)n >9 microsatellites in the bovine genome is estimated at >100, cc (T6)n >9 microsatellites in the bovine genome is estimated at >100, cc (T6)n >9 microsatellites in the bovine such bovine microsatellites cc is summarised in the specification and indexed herein (see below). cc The sequences upstream and downstream of the microsatellite sequence camplification of the corresp, microsatellite (using the program optipring). The microsatellites may be used to identify individuals, cfor parentage testing, and in the genetic mapping of economic trait cloci, or genes involved the determinism of economic trait craits esp. in cattle, to allow selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 46 BP; 18 A; 2 C; 26 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Table 7; Page 157; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene mapping, and selective breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-284684/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antibody; sperm; S19; contraception; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF60024;
   WPI; 2001-182730/18
                                                             Herr JC,
                                                                                                                     (UYVI-)
                                                                                                                                                                            23-JUL-1999;
                                                                                                                                                                                                                                    21-JUL-2000; 2000WO-US19843
                                                                                                                                                                                                                                                                                                 01-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 AGCGAGCGGGCGAGCCGAGCGAAGGGGAAAGCGCAAGAGAGAG 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34;
                                                                                                                        UNIV VIRGINIA PATENT FOUND
                                                             Norton EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                               99US-0145512
                                                                Diekman AB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 26.8; DB 13;
Pred. No. 5.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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Search completed: March 13, 2003, 21:59:12 Job time : 426 secs
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                                                                                                                                                                         Query Match
Best Local (
                                                                The present invention relates to a recombinant antibody capable of specifically binding to sperm agglutination antigen-1 (SAGA-1). The recombinant antibody has two peptide fragments of the S19 antibody and the fragments are joined together by a linker. The recombinant monoclonal antibodies are useful in a passive immunity composition for contraception as they inhibit the ability of sperm to fertilize an egg. The antibodies may be used e.g. as an active ingredient of a spermistatic agent, or as a component of a
                                                                                                                                                                                                                          Sequence 45 BP; 2 A; 10 C; 27 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                      spermicidal contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 43; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New recombinant antibody derivative of monoclonal antibody S19, useful as a reagent for purifying or detecting human spermatozoa, as an active ingredient of a spermistatic agent, or as a component of a spermicidal contraceptive
                                                                                                                                                              Y Match 1.7%; Score 26.4; DB 22; Local Similarity 75.0%; Pred. No. 6.8e+04;
                                                                                                                                                    33; Conservative
                                                                                                                                              0; Mismatches 11; Indels
                                                                                                                                                                                     Length 45;
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1: /cgn2_6/ptodata/1

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ster than or equal to the score of the result being printed,
rived by analysis of the total score distribution.
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US-08-9617-871-6
US-08-96-17-871-1
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US-08-954-698-43
US-08-954-698-43
US-08-958-026A-13
US-08-958-026A-13
US-08-958-026A-13
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US-09-9813-781-129
US-09-9813-781-13
US-08-08-747-2
US-09-9813-123-13
US-09-9813-133-19
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TITLE OF INVENTION: METHODS
FILE REFERENCE: BIV-067.01
CURRENT APPLICATION NUMBER: US/09/325,256
CURRENT FILING DATE: 199-06-03
PRIOR APPLICATION NUMBER: 60/099,800
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/078,935
PRIOR APPLICATION NUMBER: 60/078,935
PRIOR APPLICATION NUMBER: 60/078,685
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/067,423
PRIOR FILING DATE: 1998-12-03
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APPLICANT:
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SOFTWARE: PatentIn V
SEQ ID NO 31
LENGTH: 49
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                                Matches
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                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                       TYPE: DNA
                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                 FEATURE:
         703 GGTGAAAGCAGAACTCGGTGGCGGCCAAATCGGGAGGCTGCT 746
                                        Match 2.3%;
Local Similarity 88.6%;
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US-08-171-389-481
US-08-171-39-36-422
US-08-123-936-481
US-08-123-936-481
US-08-475-228A-422
US-08-475-228A-481
US-08-491-828-2
US-08-491-828-2
US-08-482-080A-422
US-08-482-080A-481
US-08-907-392A-2
US-08-908-482-747-10
US-08-983-607-42
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Pred. No.
ALIGNMENTS
                                   Mismatches
                                                      DB 4; Length 49
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; Sequence 7, Application US/08068747
; Patent No. 5695933
                                  US-08-068-747-7
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                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 08/86
PRIOR FILING DATE: 1997-06-09
PRIOR APPLICATION NUMBER: PCT/F
PRIOR FILING DATE: 1995-11-08
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
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Best Local
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TITLE OF INVENTION: Full-Length Serine Protein Kinase in Brain and Pancreas
FILE REFERENCE: 16U 101 V1
CURRENT APPLICATION NUMBER: US/09/930,181
CURRENT FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 18
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                                                                                                                                                                   Matches
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CURRENT APPLICATION NUMBER: US/09/580,923
CURRENT FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Blanche, Francis
TITLE OF INVENTION: PURIFICATION OF A TRIPLE HELIX FORMATION WITH
TITLE OF INVENTION: IMMOBILIZED OLIGONUCLEOTIDE
FILE REFERENCE: 03804.0138-01
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                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 50
                                                                                                                                                                                                                                                                                                                                    LENGTH: 50
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Similarity 73.5%;
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                                                                                                                                                                                     Score 26; DB 4;
Pred. No. 4.5e+03;
                                                                                                                                                                       Mismatches
                                                                                                                                                                          15:
                                                                                                                                                                                                        Length 50
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US-08-846-020A-6
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                                                                                                                                                                                                                                                                                    Patent No. 6090547
                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                   1351 CAGCGGCGGGGGACCGCGGGGGGGGGGGGGGG 1383
                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: CHOATE, HALL
STREET: 53 State Street
                                                                                                                                                             TITLE OF INVENTION: 5-Lipoxygenase Gene Sequence
TITLE OF INVENTION: Polymorphisms and Their Use in Classifying Patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 1; MOLECULE TYPE:
                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                       APPLICANT:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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COMPUTER: IBM PC OPERATING SYSTEM:
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                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 28; Conser
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REFERENCE/DOCKET NUMBER: MIT-6141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DESCRIPTION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                          02109-2891
                                                                                                   Boston
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                                                                          USA
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                                                                                                                                                                                                                                                         Drazen M.D., Jeffrey M.
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Housman, David E.
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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/desc = "Synthetic"
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Pred. No. 6.6e+03;
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TELEFAX: (617) 248 400 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
               NAME: Jarrell Ph.D., Brenda H.
REGISTRATION NUMBER: 39,223
REFERENCE/DOCKET NUMBER: 0926
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-5000
TELEPAX: (617) 248 4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC_DOS_MS-DOS SOFTWARE: Patentin Rclease #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: 5-Lipoxygenase Gene Sequence
TITLE OF INVENTION: Polymorphisms and Their Use in Classifying Patients
                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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NAME: Jarrell Ph.D. Brenda H.
REGISTRATION NUMBER: 39, 223
REFERENCE/DOCKET NUMBER: 00926(
TELECOMMUNICATION INFORMATION:
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                                                                                                                                    FILING DATE:
                                                                                                                                                                                   CLASSIFICATION:
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Local Similarity 76.9%;
                                                                                                       NAME:
                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/617,871
                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                    CITY: Boston
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Pred. No. 8.3e+03;
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STRANDEDNESS:
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; TOPOLOGY: line;
; MOLECULE TYPE: cl
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US-08-748-591-11
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US-08-356-060A-43
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                                                                                                                                  Matches
                                                                                                                                                Best Local Similarity
                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 854-0875
INFORMATION FOR SEQ ID NO:
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TOPOLOGY: not relevant
MOLECULE TYPE: DNA (genomic)
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LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                              NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/748,591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Epstein, Ervin
APPLICANT: Hu, Zhilan
APPLICANT: Bonifas, Jeanette
TITLE OF INVENTION: Mutant Human Hedgehog Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
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                                                                                                  524 ACCGAGGGCTGGGACGAAGATGGC 547
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CITY: Menlo Park
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Local Similarity 76.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                  TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                LENGTH:
                                                                 1 ACCGAGGGCTGGGACGAAGATGGC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30;
                                                                                                                                  24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94025
                                                                                                                                                                                                                                                             nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fish and Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                    CDNA
                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                  435
                                                                                                                                                1.5%; Score 24; DB 1; I
100.0%; Pred. No. 9.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                    06510/067001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24.6; DB 4;
Pred. No. 8.3e+03;
                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 4;
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                                                                                                                                                                   Length 24;
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                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                               ; Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 43, Application US/08356060A Patent No. 5844079
                                                                                                                                                                                                                                                                                             Sequence 43, Application US/08460900C Patent No. 6165747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ. ID NO: 43:
                                                                                                                                                                                                                                                                              GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Vertebrate Embryonic Pattern-Inducing TITLE OF INVENTION: Proteins and Uses Related Thereto NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: McMahon, Andrew P. APPLICANT: Tabin, Clifford J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 30-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                              APPLICANT: Ingham, Phillip W.
APPLICANT: McMahon, Andrew P.
APPLICANT: Tabin, Clifford J.
APPLICANT: Bumcrot, David A.
APPLICANT: Marti-Gorostiza, Elisa
APPLICANT: Marti-Gorostiza, Elisa
TITLE OF INVENTION: Peroteins and Uses Related Thereto
                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                          524 ACCGAGGGCTGGGACGAAGATGGC 547
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/176,427 FILING DATE: 30-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/356,060A FILING DATE: 14-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 1.5%; Score 24; DB 2; Local Similarity 100.0%; Pred. No. 9.9e+03 es 24; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
COUNTRY: UZIP: 02109
                                  STATE:
                                                    CITY: Boston
                                                                     ADDRESSEE: FOLEY, HOAG & ELIOT STREET: One Post Office Square
                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02109
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                                    MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 base pairs
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                   USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                            HOAG & ELIOT LLP
                                                                                                                                   62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 24
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COMPUTER READABLE FORM:

MEDIUM TYPE:

Floppy disk

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 43, Application US/08674509B
Patent No. 6261786
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ingham, Phillip W. APPLICANT: McMahon, Andrew P. APPLICANT: Tabin, Clifford J. APPLICANT: Marigo, Valeria TITLE OF INVENTION: SCREENING A TITLE OF INVENTION: AND ANTAGON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/460,900
FILING DATE: 05-JUN-1995
                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY, HOAG & ELIOT LLP
STREET: One Post Office Square
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: 1i
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            524 ACCGAGGGCTGGGACGAAGATGGC 547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 30-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 5-JUNE-19
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APPLICATION NUMBER: US 08/356,060
FILING DATE: 14-DEC-1994
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                                                           CLASSIFICATION:
                                                                          FILING DATE:
                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/435,093 FILING DATE: 4-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
                                                                                                                                                                                                                   02109-2170
                                                                                                                                                                                                                                                                     Boston
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                                                                                                                                                                                                                                                    MA
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                                                                                                                                                                                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (617) 832-7000
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                                                                       02-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JMBER: US 08/176,427
30-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JMBER: US/08/460,900C
5-JUNE-1995
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                                                                                                                                                                                                                                                                                                                                         AND ANTAGONISTS
48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.5%; Score 24;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                      SCREENING ASSAYS FOR HEDGEGOG AGONISTS
                                                                                       US/08/674,509E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HMV-006.05
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9.9e+03;
hes 0; Indels
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RESULT 11
US-08-954-698-43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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             TELEFAX: 617-832-7000 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: US/08/954,698 FILING DATE: 20-OCT-1997 PRIOR APPLICATION NUMBER: US 08/462,386 FILING DATE: 05-JUN-1995 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/462,386 FILING DATE: 05-JUN-1995 PRIOR APPLICATION DATA:
                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 30-DEC-1993
                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/435,093
FILING DATE: 04-MAY-1995
PRIOR APPLICATION DATA:
                                                                      TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ingham, Phillip W.
APPLICANT: McMahon, Andrew P.
APPLICANT: Tabin, Clifford J.
TITLE OF INVENTION: Vertebrate Embryonic Pattern-Induc
TITLE OF INVENTION: Proteins and Uses Related Thereto
SEQUENCE CHARACTERISTICS:
                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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REFERENCE/DOCKET NUMBER: HM
TELECOMMUNICATION INFORMATION:
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                                                                                                       NAME: Vincent, Matthew P. REGISTRATION NUMBER: 36,7
                                                    TELEPHONE:
                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/356,060 FILING DATE: 14-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 02109-2170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08954698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: FOLEY, HOAG & ELIOT LLP
One Post Office Square
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                    617-832-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          617-832-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other nucleic acid
/desc = "primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                                                  US 08/176,427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.5%; Score 24; DB 4; Length 24; 100.0%; Pred. No. 9.9e+03;
                                                                                    36,709
ER: HMV-006.10
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                                                                                                                                             US-08-957-874-43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                     Query Match
          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 5-JUNE-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/435,093
FILING DATE: 4-MAY-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/356,060
FILING DATE: 14-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION 
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                                                                                                                                                                                                                                                                                                                              TELEPHONE: (617) 832-1000
TELEFAX: (617) 832-7000
INFORMATION FOR SEO ID NO: 43:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Vertebrate Employment of Invention: Proteins and Uses Related Thereto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ingham, Phillip W. APPLICANT: McMahon, Andrew P. APPLICANT: Tabin, Clifford J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: HMT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: ASCII(text)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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                                                                                                                                                                         TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: STREET: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       524 ACCGAGGGCTGGGACGAAGATGGC 547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 5-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0. FILING DATE: 20-OCT-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
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                                      Local Similarity
                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                   TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                         LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vincent, Matthew P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: FOLEY, HOAG & ELIOT LLP
One Post Office Square
                                                                                                                                                                                                                                                                                                             24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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100.08; F1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vertebrate Embryonic Pattern-Inducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.5%; Score 24;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/08/957,874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US 08/462,386
   1.5%; Score 24; DB 4; Length 24; L00.0%; Pred. No. 9.9e+03; Lye 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HMV-006.09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4; Length 24; 9.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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US-08-258-026A-13
                                                                                                                                                                                                               US-09-813-781-129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13, Application US/08258026A Patent No. 5516637
                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                          Sequence 129, Application US/09813781 Patent No. 6405989
                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEPAX: (617) 523-6400
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 13:
APPLICANT: WEIDANZ, JON A.
APPLICANT: CARD, KIMBERLYN F.
APPLICANT: WONG, HING C.
APPLICANT: WONG, HING C.
TITLE OF INVENTION: FUSION PROTEINS COMPRISING BACTERIOPHAGE COAT PROTEIN
TITLE OF INVENTION: AND A SINGLE-CHAIN T-CELL RECEPTOR
FILE REPERENCE: 46745(1758)
CURRENT APPLICATION NUMBER: US/09/813,781
CURRENT FILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 130
                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
TYPE: nucleic acid
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TITLE OF INVENTION: A METHOD FOR DISPLAYING PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      524 ACCGAGGGCTGGGACGAAGATGGC 547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 10-JUN CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 342
REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME:
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                                                                                                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Resnick, David R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MASSACHUSETTS
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                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   David G. Conlin; DIKE, BRONSTEIN, ROBERTS & CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              unknown
                                                                                                                                                                                                                                                                                                                                                                         1.5%;
75.0%;
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                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                         Score 24; DB 1;
Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 45;
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                                                                Matches
                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13, Application:
                                                                                                                                                                                        TELEFAX: (617) 523-6400
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. SEQ ID NO 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 1.5%;
Best Local Similarity 75.0%;
               TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 45
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: A METHOD TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                 STRANDEDNESS:
TOPOLOGY: un
                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/258,026 FILING DATE: 10-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY:
STATE:
                                                                                                                                                                                                                                                                                       NAME: Resnick, David R
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 42838
                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US95/07541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER:
2 GTGGCGGTGGCAGCGGCGGTGGTTCCGGAGGCGGCGG 41
                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02109
                                                                                                                                                          nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MASSACHUSETTS
                                                                                                                                                                        45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application PC/TUS9507541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: David G. Conlin; DIKE, BRONSTEIN, E: ROBERTS & CUSHMAN 130 WATER STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rhode, Peter R.
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SD
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                                                                                                                                  unknown
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VENTION: A METHOD FOR DISPLAYING
                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stinson,
                                                                                                                                              unknown
                                                                         1.5%;
75.0%;
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                                                                                                                                                                                                                                                                                        34235
                                                                                     Score 24; DB 5;
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                                                                         Pred. No. 1.1e+04;
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Pred. No.
                                                            Mismatches
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                                                         10;
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RESULT 17
US-09-499-884-12
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                           NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn v
SEQ ID NO 12
                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: St. Clair, Daret
                                                                                                                                                                                                               Sequence 12, Application US/09499884
Patent No. 6265172
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                                                                                                             APPLICANT: St. Clair, Daret
APPLICANT: Urano, Muneyasu
APPLICANT: Kasarskis, Edward
TITLE OF INVENTION: DIAGNOSTIC TEST ANI
TITLE OF INVENTION: ASSOCIATED DISEASES
                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                   CURRENT APPLICATION NUMBER: US/09/499,884
CURRENT FILING DATE: 2000-02-08
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                                                                                                        FILE REFERENCE:
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Synthetic"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 617-861-6240
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MEDIUM TYPE: Floppy
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APPLICANT: Hudson, Thomas J.
APPLICANT: Housman, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                         30 CGGCGGCGGCGGCGGCGGCGGCGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 20
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                       Local Similarity es 26; Conserv
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REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
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                38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
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                                        PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Massachusett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30 base pairs
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                                                                                                                                                                                                                                                                                                                                                                    Conservative
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IBM PC compatible
                                                                                                       50229-180
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28-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single
                                                                                                                                                                                                                                                                                                                                                                                   1.5%;
                                                                                                                            DIAGNOSTIC TEST AND THERAPY FOR MANGANESE SUPEROXIDE DISMU
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ORGANISM: Homo sapiens

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; LOCATION: (40)..(48)
; OTHER INFORMATION: n = a,
US-09-497-933A-19
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           δõ
                                                                                                            US-09-930-181-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                     Sequence 13, Application US/09930181 Patent No. 6455292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wagner, Robert Jr. E.
TITLE OF INVENTION: METHODS FOR DETECTION OF A TRIPLET REPEAT BLOCK AND A
TITLE OF INVENTION: FUNCTIONAL MISMATCH BINDING PROTEIN IN A BIOLOGICAL
TITLE OF INVENTION: FLUID SAMPLE
                                                                                                                                                                            SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 9408-044
CURRENT APPLICATION NUMBER: US/09/497,933A
CURRENT FILING DATE: 2000-02-04
                                             Matches
                                                                            Query Match
                                                                                                                                                                                                     FILE REFERENCE: 16U 101 V1
CURRENT APPLICATION NUMBER: US/09/930,181
CURRENT FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                   APPLICANT: OriGene Technologies
TITLE OF INVENTION: Full-Length Serine Protein Kinase in Brain and Pancreas
FILE REFERENCE: 16U 101 V1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: modified_base
LOCATION: (1)..(9)
OTHER INFORMATION: n = a, c, g or t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 48
                                                                                                                                                         LENGTH: 50
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                                                              Local Similarity
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32;
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Similarity 76.3%;
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                                             Conservative
                                                              1.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 23.6;
                                                           Score 23.6; DB 4; Length 50; Pred. No. 1.3e+04;
                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.3e+04;
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                                                  Indels
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RESULT 20
US-08-748-591-12/c
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                                                                                                                                                                                                                                                                                                                            RESULT 21
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                                                                                                                                                                                                                                                     ; GENERAL INFORMATION:
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SEQ ID NO 5
                                                                                FILE REFERENCE: sequence listing CURRENT APPLICATION NUMBER: US/09/083,123 CURRENT FILING DATE: 1998-05-22 EARLIER APPLICATION NUMBER: EP 95118600.6 EARLIER FILING DATE: 1995-11-25
                                                                                                                                                                             APPLICANT: Orum, Hendrik
APPLICANT: Seeger, Corina
TITLE OF INVENTION: Method for Generating Multiple Double Stranded Nucleic
TITLE OF INVENTION: Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (415) 854-0875
INFORMATION FOR SEQ ID NO:
                            EARLIER APPLICATION NUMBER: PCT/EP96/05149
EARLIER FILING DATE: 1996-11-22
NUMBER OF SEQ ID NOS: 8
               SOFTWARE: PatentIn Ver. 2.0
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APPLICANT: Hu, Zhilan
APPLICANT: Hu, Zhilan
APPLICANT: Bonifas, Jeanette
TITLE OF INVENTION: Mutant Human Hedgehog Gene
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                616 CAGCAAGTACGGCATGCTGGCCCGC 640
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/748,591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER:
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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US-08-340-045-7
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                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-325-256-30
                                                                                                                                  RESULT 23
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SOFTWARE: Par
SEQ ID NO 30
SEQTH: 47
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                                                                                    Sequence 7, Patent No. 5
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                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                               Query Match
                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-12-03
PRIOR APPLICATION NUMBER: PCT,
PRIOR FILING DATE: 1998-12-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/325,256
CURRENT FILING DATE: 1999-06-03
PRIOR APPLICATION NUMBER: 60/099,800
PRIOR APPLICATION NUMBER: 60/099,800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: HYDROPHO
TITLE OF INVENTION: METHODS
FILE REFERENCE: BUY-067.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
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PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
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   APPLICANT:
                              APPLICANT:
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                                                          APPLICANT:
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                APPLICANT:
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                                                                                       , Application US/08340045 5637495
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GALDES,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PEPINSKY, R. BLAKE
OPPENHEIM, AMOS VOGEL, TIKVA
                             GORECKI, MARIAN
LEVANON, AVIGDOR
                                                        AVIV, HAIM
                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                              100.0%;
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100.0%; Pred. No.
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Pred. No. 2.1e+04;
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. 2.7e+04;
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US-08-871-302A-7
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REGISTRATION NUMBER: 28,678

REFERENCE/DOCKET NUMBER: 2006

TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
TELEPAX: 212-391-0525
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                  APPLICANT: ZEELON, ELISHA
APPLICANT: ZEEVIN, MENACHEM
TITLE OF INVENTION: PLASMIDS FOR PRODUCTION OF HUMAN GROWTH
TITLE OF INVENTION: HORMONE OR POLYPEPTIDE ANALOG THEREOF, HOSTS CONTAINING
TITLE OF INVENTION: METHODS
TITLE OF INVENTION: METHODS
TITLE OF INVENTION: METHODS
TITLE OF INVENTION: METHODS
NUMBER OF SEQUENCES: 32
                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                         APPLICANT:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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HYPOTHETICAL: NO
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SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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APPLICANT:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       789 AGCTGGTGAAGGACCTGAGCCCCGGGGACCGCG 821
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                STATE:
                                   CITY: NEW YORK
                                                                   ADDRESSEE:
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                     3 AGCTGGTGCAGGTGCTGAGCCCGGAGCACAGCG
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STRANDEDNESS: unkn
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                NEW YORK
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                                              E: COOPER & DUNHAM LLP
1185 AVENUE OF THE AMERICAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NEW YORK
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1185 AVENUE OF THE AMERICAS
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                                                                                                                                                                                                                                                             LEVANON, AVIGDOR
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ZEEVI, MENACHEM
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SYSTEM: PC-DOS/MS-DOS
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HORMONE OR POLYPEPTIDE ANALOG THEREOF, HOSTS CONTAINING
THE PLASMIDS, PRODUCTS MANUFACTURED THEREBY, AND RELATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/08/340,045
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Pred. No. 2.8e+04;
^ "'smatches 7;
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CURRENT APPLICATION NUMBER: US/09/485,737B
CURRENT APPLICATION NUMBER: COT/EP 98/05165
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1997-08-18
NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
NUMBER: OF SEQ ID NOS: 104
SOFTWARE: PATENTIN VETSION 3.0
SEQ ID NO 5
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                                                                                                                             ; OTHER INFORMATION: GENOMIC US-09-485-737B-5
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                                      Query Match
Best Local Similarity 75.0
Matches 27; Conservative
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TELEPHONE: 212-278-0400
TELEPAX: 212-391-0525
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Buyse, Marie-Ange
APPLICANT: Sablon, Erwin
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 35 base pairs
TYPPE: nucleic acid
STRANDEDNESS: unknown
TOPOLOGY: linear
MOLECULE TYPE: DNA (genom
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                         TYPE: DNA
ORGANISM: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: WHITE, JOHN P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 20065-G/JPW/GJG
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL:
ANTI-SENSE:
                                                                                                                                                                                                                                         LENGTH: 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
757 GGCCACGGTGCACCTGGAGCAGGGCGGCACCAAGCT 792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 AGCTGGTGCAGGTGCTGAGCCCGGAGCACAGCG 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/871,302A
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                                                                1.4%;
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                                             0; Mismatches
                                                                  Score 21.6; DB 4; Length 40; Pred. No. 3.1e+04;
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Search completed: March 13, 2003, 23:59:15 Job time: 96 secs

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Perfect score:
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and is derived
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Bd
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                       d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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seq length:
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Match
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/cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
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US-10-135-984-7

US-10-135-984-7

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US-09-992-598-122

US-09-998-735-122

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| US-09-997-748-313
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Sequence 20, Appl Sequence 19, Appl Sequence 1, Appli Sequence 7, Appli Sequence 7, Appli Sequence 7, Appli Sequence 269, App Sequence 269, Appli Sequence 29, Appli Sequence 29, Appli Sequence 30, Appli Sequence 30, Appli Sequence 31, Appli		Sequence 3, Appli Sequence 8, Appli Sequence 8, Appli Sequence 10, Appli Sequence 37, Appl Sequence 37, Appl Sequence 37, Appl Sequence 37, Appl Sequence 27, Appl Sequence 4, Appli Sequence 4, Appli	Sequence 53, Appl Sequence 53, Appl	sequence 122, Sequence 122, Sequence 122, Sequence 122, Sequence 122, Sequence 123, Sequence 1131, Sequence 1711, Sequence 53, All equence 53, All equence 53, All equence 53, All	113, A 113, A 222, A 1122, A 1
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Patent No. US20020019350A1
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982
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         Query Match
Best Local Similarity
 Matches
                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/782,650 CURRENT FILING DATE: 2001-02-12 PRIOR APPLICATION NUMBER: US 09/324,079 PRIOR FILING DATE: 1999-06-01 PRIOR APPLICATION NUMBER: US 09/327,045 PRIOR FILING DATE: 1999-06-07
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                                                                                                                PRIOR APPLICATION NUMBER: PCT/US00/14988
PRIOR FILING DATE: 2000-05-31
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                            APPLICANT: Edwards LifeSciences Corporation TITLE OF INVENTION: Targeted Angiogenesis FILE REFERENCE: 20553D-00061US
                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                       APPLICANT: Levine, Arnold J. APPLICANT: Mitterer, Artur
                                                                                                                                                                                                                                            APPLICANT:
                                             OTHER INFORMATION: Description of Artificial Sequence:annealed OTHER INFORMATION: oligonucleotide
                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
                                                                  FEATURE:
                                                                                              ENGTH: 50
                                                                                                         ID NO 16
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Falkner, Falko-Guenter
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Dorner, Friedrich
Conservative
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85.4%;
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US-10-056-414-577
US-09-927-122-49
US-09-9504-231A-2464
US-09-544-231A-2664
US-09-274-553D-2453
US-09-814-924-2
US-10-116-288-10
US-09-918-696-43
US-09-918-696-43
US-09-918-696-44
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0 US-09-778-175-16
0 US-09-335-218-16
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0 US-09-179-536B-245
US-09-864-785-742
US-09-864-785-742
US-09-874-547-75
0 US-09-870-203A-23
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0 US-09-9870-203A-23
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US-09-940-550A-7
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          Score 31.4; DB 10;
Pred. No. 1.2e+03;
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 Mismatches
                    Length 50;
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Sequence 7, Appli
Sequence 91, Appli
Sequence 27, Appl
Sequence 2453, Ap
Sequence 2664, Appl
Sequence 2664, Appli
Sequence 2, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 11, Appli
Sequence 16, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 11, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 1133, Ap
Sequence 72, Appl
Sequence 73, Appl
Sequence 75, Appl
Sequence 75, Appl
Sequence 23, Appl
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24, Appl
85, Appl
   Gaps
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; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: Oligonucleotide US-09-883-848A-37
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US-09-883-848A-37/c
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                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic RNA US-09-828-034-7
                                                                                                                                                                                                                                                                APPLICANT: Hong, Zhi
APPLICANT: Hong, Zhi
APPLICANT: Ferrari, Eric
FITTLE OF INVENTION: HCV REPLICASE COMPLEXES
FILE REFERENCE: IN01165
CURRENT APPLICATION NUMBER: US/09/828,034
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: U.S. 60/195,852
PRIOR APPLICATION NUMBER: U.S. 60/195,852
PRIOR APPLICATION NUMBER: 0.50/195,852
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
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LENGTH: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL
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Patent No. US20020064771A1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ling, L.
APPLICANT: Sanicola-Nadel, M.
APPLICANT: Sanicola-Nadel, M.
TITLE OF INVENTION: ANGIOGENESIS-MODULATING COMPOSITIONS AND USES
FILE REFERENCE: CIBT-P01-119
CURRENT APPLICATION NUMBER: US/09/883,848A
CURRENT FILING DATE: 2001-06-18
CURRENT FILING DATE: 2001-06-18
                                                                                                                                                                                                                                                   SEQ ID NO 7
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Best Local
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PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 48
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                                                                                                                                                                                                                LENGTH: 30
TYPE: RNA
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                                   1546 GGGGGCCGGGGAGGGGGC 1575
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INFORMATION
                                                                         25;
                                                                                         Similarity
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                                                                         Conservative
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                                                                                         1.4%;
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Pred. No.
                                                                         0;
                                                                                         Score 22;
Pred. No.
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                                                                                             6.2e+04;
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                                                                                                            DB 10;
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US-10-006-856A-151

RESULT 4

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                                                                                                                                               ; TYPE: DNA; ORGANISM: adeno-associated virus US-10-135-984-6
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US-10-006-856A-151
                                                                            Matches
                                                                                                           Query Match
                                                                                                                                                                                                                    SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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Publication No. US20020182595A1
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Best Local :
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CURRENT FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: 60/286951
PRIOR FILING DATE: 2001-04-27
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Anton J. Cathomen
TITLE OF INVENTION: METHOD OF IDENTIFYING CELLULAR
TITLE OF INVENTION: REGULATORS OF ADENO-ASSOCIATED VIRUS (AAV)
FILE REFERENCE: SALKINS.041A
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                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Matthew D. Weitzman APPLICANT: Anton J. Cathomen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P28309D1C14
CURRENT APPLICATION NUMBER: US/10/006,856A
CURRENT FILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 477
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                   LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       763 GGTGCACCTGGAGCAGGGCGGCACCAAGCTGGTGAAGGA 801
                                                                                           Local Similarity
5 GTGAGCGAGCGAGCGCAGGTGAGCGAGCGAGCGCGCAG 44
                                   1 GCGAGGCAGCCAGCGAGGGAGAGAGCGAGCGGGCGAGCCG 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                   45.
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Godowski, Paul J.
Grimaldi, Christopher J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fong,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Paoni, Nicholas F.
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Hillan, Kenneth J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gao,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eaton, Dan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Botstein, David
                                                                          Conservative
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                                                                                         1.3%;
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                                                                                       Score 20.8; DB 9;
Pred. No. 1.1e+05;
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Pred. No. 8.3e+04;
                                                                      Mismatches
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                                                                                                      DB 9;
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                                                                                                        Length 45;
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RESULT 6

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US-10-135-984-7/c; Sequence 7, Application US/10135984; Publication No. US20020182595A1
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                                                                                                                                                                                                                                             RESULT 8
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                                                                                                                                                      GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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APPLICANT: Anton J. Cathomen
TITLE OF INVENTION: METHOD OF IDENTIFYING CELLULAR
TITLE OF INVENTION: REGULATORS OF ADENO-ASSOCIATED VIRUS (AAV)
FILE REFERENCE: SALKINS.041A
CURRENT APPLICATION NUMBER: US/10/135,984
CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: REGULATORS OF ADENO-ASSOCIATED VIRUS (AAV) FILE REFERENCE: SALKINS.041A CURRENT APPLICATION NUMBER: US/10/135,984 CURRENT FILING DATE: 2002-08-05 PRIOR APPLICATION NUMBER: 60/286951 PRIOR FILING DATE: 2001-04-27
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                                                                                                           APPLICANT: Matthew D. Weitzman APPLICANT: Anton J. Cathomen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Anton J. Cathomen
TITLE OF INVENTION: METHOD OF IDENTIFYING CELLULAR
TITLE OF INVENTION: REGULATORS OF ADENO-ASSOCIATES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Matthew D. Weitzman APPLICANT: Anton J. Cathomen
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TITLE OF INVENTION: METHOD OF IDENTIFYING CELLULAR
TITLE OF INVENTION: REGULATORS OF ADENO-ASSOCIATED VIRUS (AAV)
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CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: SALKINS.041A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: adeno-associated virus
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                                                                                                                                                                                                                                                                                                        45 GTGAGCGAGCGAGCGCGCGCGCGCGCGCGCGCGCAG
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0; Mismatches
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Pred. No. 1.1e+05;
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: LCATION: (0) ...(0)
: OTHER INFORMATION: phosphorothioate backbone US-09-888-326-172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Methods for Enhancing Antibody-Induced TITLE OF INVENTION: Cell Lysis and Treating Cancer FILE REFERENCE: C1039/7052 (AWS)
CURRENT APPLICATION NUMBER: US/09/888,326
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: US 60/213,346
PRIOR APPLICATION NUMBER: US 60/213,346
PRIOR FILING DATE: 2000-66-22
NUMBER OF SEQ ID NOS: 848
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 172
                                                                                                                                          GENERAL INFORMATION:
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            CURRENT APPLICATION NUMBER: US/10/027,961A
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: 09/599,781
PRIOR FILING DATE: 2000-06-21
PRIOR APPLICATION NUMBER: 08/930,589
PRIOR FILING DATE: 1996-02-22
PRIOR APPLICATION NUMBER: PCT/CA94/00501
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PRIOR FILING DATE: 2001-04-27
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                                                                                                                                   FILE REFERENCE: 19029PCADA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
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FILING DATE: 1999-09-13
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Pred. No. 1.1e+05;
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Pred. No. 1.2e+05;
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US-10-072-152-32
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                                                                                                                                                    ; GENERAL INFORMATION:
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                                                                                                                                                                  Patent No. US20020142438A1
                                                                                                                                                                                  Sequence 32, Application US/10072152
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                                                                                                                                   Andersen, Lene N.
               Kauppinen, Markus
Schnorr, Kirk
                                               Moller, Soren Glad, Sanne O.
                                                                                Bjornvad, Mads
                                                                                            Lange, Niels E.
 Kongsbak, Lars
                                                                                                                 Schulein, Martin
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GENERAL INFORMATION:
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Patent No. US20020001798A1
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PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MORROW, PHILLIP
TITLE OF INVENTION: NEUTRALIZING HIGH AFFINITY HUMAN MONOCLONAL ANTIBODIES
TITLE OF INVENTION: SPECIFIC TO RSV F-PROTEIN AND METHODS FOR THEIR
TITLE OF INVENTION: MANUFACTURE AND THERAPEUTIC USE THEREOF
FILE REFERENCE: 037003-0275759
CURRENT APPLICATION NUMBER: US/09/740,002
CURRENT FILING DATE: 2000-12-20
CURRENT FILING DATE: 2000-12-20
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PRIOR FILING DATE: 1993-09-27
PRIOR APPLICATION NUMBER: 08/064,271
PRIOR FILING DATE: 1993-05-06
PRIOR APPLICATION NUMBER: 07/994,760
PRIOR FILING DATE: 1992-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 09/335,697 PRIOR FILING DATE: 1999-06-18
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                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     183 TCCTCGTCTCCTCGCTGCTGGTATGCTCGGGACTGGCG 220
                                            19 GAGAGAGCGAGCGGGCGAGCGAGGGAAGGGAAA 56
                                                                                                                  Local Similarity
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2 GAGAGAGAGAGAGAGCCAGATACACGATGTGAAA 39
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71.1%;
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                                                                                                                  Score 20.4; DB 10
Pred. No. 1.3e+05;
                                                                                              Mismatches
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US-09-992-598-122; Sequence 122, A
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APPLICANT: Ashkenaz
APPLICANT: Baker, K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 32
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                                                       APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C20
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: APPLICANT:
CURRENT APPLICATION NUMBER: US/09/92,598
CURRENT EILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
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PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/067,249
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PRIOR FILING DATE: 1997-11-24
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PRIOR FILING DATE: 1998-11-24
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CURRENT FILING DATE: 2002-02-07
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FILING DATE: 1998-11-02
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Baker, Kevin P.
                                                                                                                         Watanabe, Colin K. Williams, P. Mickey Wood, William I.
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Stewart, Timothy A.
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Godowski, Paul J.
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                                                                                                                                                                         Daniel
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PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12 PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: 60/075945 RIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR FILING DATE: 1997-11-13 PRIOR APPLICATION NUMBER: 60/065311 PRIOR APPLICATION NUMBER: 60/062250 APPLICATION NUMBER: FILING DATE: 1998-06 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: 1998-0 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: FILING DATE: FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/0 TILING DATE: 1998-06-10 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/088742 FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/088202 FILING DATE: 1998-06-05 APPLICATION NUMBER: FILING DATE: 1998-06-05 APPLICATION NUMBER: APPLICATION NUMBER: 60/088028 FILING DATE: 1998-06-04 FILING DATE: 1998-06-05 APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: FILING DATE: FILING DATE: 1998-01 APPLICATION NUMBER: APPLICATION NUMBER: 60/087759 FILING DATE: 1998-06-02 FILING DATE: FILING DATE: 1998-05-28 APPLICATION NUMBER: 60/087607 FILING DATE: FILING DATE: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/087106 APPLICATION NUMBER: APPLICATION NUMBER: 60/083322 FILING DATE: 1998-04-28 FILING DATE: 1998-02-25 APPLICATION NUMBER: 60/078910 FILING DATE: FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/087827 FILING DATE: APPLICATION NUMBER: 60/066770 1998-06-1998-06-1 1998-06-11 1998-06-10 1998-06-1998-06-10 1998-06-10 1998-06-10 1998-06-10 1998-06-1998-06-04 1998-06-04 1998-06-05 1998-06-04 1998-06-04 1998-06-04 1998-06-03 1998-06-04 1998-06-02 1998-06-02 1998-05-07 1998-03-20 1997-11-24 .998-06-04 60/089105 60/088876 60/088858 60/088826 60/088655 60/088217 60/088861 60/088824 60/088810 60/088738 60/088734 60/088212 60/088167 60/088326 60/088033 60/088030 60/088029 60/088026 60/088025 60/088021 60/087609 60/084600

FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089512 FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089514

1998-06-16

1998-06-17

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RESULT 14
US-09-989-293A-122
; Sequence 122, Application US/09989293A
; Patent No. US20020177164A1
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          PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-13
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C66
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FILING DATE: 1998-07-02
APPLICATION NUMBER: 60/091978
FILING DATE: 1998-07-07
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FILING DATE: 1998-07-01
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APPLICATION NUMBER: 60/066770
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Williams, P. Mickey
Wood, William I.
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Roy, Margaret Ann
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Pred. No.
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APPLICATION NUMBER: 60/090676 FILING DATE: 1998-06-25 APPLICATION NUMBER: 60/090678 FILING DATE: 1998-06-25

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PRIOR PRIOR

FILING DATE: 1998-06-25 APPLICATION NUMBER: 60/090694 FILING DATE: 1998-06-25

APPLICATION NUMBER: 60/090690

R APPLICATION NUMBER: 60/090695
R FILING DATE: 1998-06-25
R APPLICATION NUMBER: 60/090696
R FILING DATE: 1998-06-25
R APPLICATION NUMBER: 60/090862
R FILING DATE: 1998-06-26

PRIOR PRIOR

FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-24

60/090444

APPLICATION NUMBER: 60/090445

1998-06-24

60/090472 60/090535

APPLICATION NUMBER: 60/090435 APPLICATION NUMBER: 60/090431 FILING DATE: 1998-06-24

FILING DATE: 1998-06-24 APPLICATION NUMBER:

60/090429

APPLICATION NUMBER: 60/090252 FILING DATE: 1998-06-22

APPLICATION NUMBER: 60/090246

1998-06-22 1998-06-19

FILING DATE: FILING DATE:

FILING DATE: 1998-06-22 APPLICATION NUMBER: 60/090349 APPLICATION NUMBER: 60/090254

APPLICATION NUMBER: 60/090355 FILING DATE: 1998-06-23 FILING DATE: 1998-06APPLICATION NUMBER: 60/089948 FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-19

APPLICATION NUMBER:

60/089952

FILING DATE: 1998-06-18
APPLICATION NUMBER: 60/089908
FILING DATE: 1998-06-18

60/089947

FILING DATE: 1998-0 APPLICATION NUMBER:

APPLICATION NUMBER: 60/089653 FILING DATE: 1998-06-17 FILING DATE: 1998-06-17
APPLICATION NIMMARY APPLICATION NUMBER: 60/089599 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089598 FILING DATE: 1998-06-17 FILING DATE: 1998-06-17 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-06

APPLICATION NUMBER: 60/089801

1998-06-18

60/089907

PRIOR

APPLICATION NUMBER: 60/090542 FILING DATE: 1998-06-24

APPLICATION NUMBER: 60/090540 FILING DATE: 1998-06-24 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/ APPLICATION NUMBER: FILING DATE:

APPLICATION NUMBER: 60/ FILING DATE: 1998-06-24

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OR FILING DATE: 1998-06-04
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OR APPLICATION NUMBER: 60/088030
OR FILING DATE: 1998-06-04 DR APPLICATION NUMBER: 60/088033

R FILLING DATE: 1998-06-04

DR APPLICATION NUMBER: 60/088326

DR FILLING DATE: 1998-06-04

DR APPLICATION NUMBER: 60/088167

DR FILLING DATE: 1998-06-05 APPLICATION FILING DATE: FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: 60/088734 FILING DATE: 1998-06-10 FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-05 FILING DATE: APPLICATION NUMBER: FILING DATE: 1998-06-FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/1 FILING DATE: 1998-06-05 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/087609 FILING DATE: 1998-05-28 APPLICATION NUMBER: 60/087607 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-05
APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/083322 FILING DATE: 1998-03-20 APPLICATION NUMBER: FILING DATE: 1997-11-24 APPLICATION NUMBER: 60/ FILING DATE: 1998-02-25 NUMBER: 60/ : 1998-06-17 1998-06-12 1998-06-11 1998-06-11 1998-06-10 1998-06-10 1998-06-10 1998-06-10 1998-06-09 1998-06-10 1998-06-02 1998-06-02 1998-05-07 1998-04-28 60/089532 60/089440 60/089514 60/089512 60/088876 60/088861 60/088858 60/089105 60/088738 60/088826 60/088824 60/088810 60/088742 60/088655 60/088217 60/088212 60/088202 60/087827 60/087759 60/087106 60/078910 60/075945

PRIOR PR FILING DATE: 1998-06-26

PR APPLICATION NUMBER: 60/090863

PR FILING DATE: 1998-06-26

PR APPLICATION NUMBER: 60/091360

PR FILING DATE: 1998-07-01

PR APPLICATION NUMBER: 60/091478

PR FILING DATE: 1998-07-02 DR FILING DATE: 1998-06
RAPPLICATION NUMBER:
DR FILING DATE: 1998-06
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PRIOR APPLICATION UNMBER: 60/091519
PRIOR APPLICATION UNMBER: 60/091626
PRIOR APPLICATION UNMBER: 60/091626
PRIOR APPLICATION UNMBER: 60/091636
PRIOR FILING DATE: 1998-07-02
PRIOR FILING DATE: 1998-07-02
PRIOR APPLICATION NUMBER: 60/091978
PRIOR APPLICATION NUMBER: 60/091978
PRIOR APPLICATION NUMBER: 60/091982
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CURRENT FILING DATE: 2001-11-19
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FILING DATE: 1998-07-07
APPLICATION NUMBER: 60/092182
FILING DATE: 1998-07-09
FILING DATE: 1998-03-20
APPLICATION NUMBER: 60/083322
FILING DATE: 1998-04-28
                                                                                                         FILING DATE: 1997-11-12
APPLICATION UNMBER: 60/065311
FILING DATE: 1997-11-13
APPLICATION NUMBER: 60/066770
FILING DATE: 1997-11-24
FILING DATE: 1997-11-24
                                                     APPLICATION NUMBER: 60/075945 FILING DATE: 1998-02-25 APPLICATION NUMBER: 60/078910
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Williams, P. Mickey
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Pred. No. 1.5e+05;
0; Mismatches 15; Indels
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PRIOR PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: 60/084600 PRIOR FILING DATE: 1998-05-07 PRIOR FILING DATE: 1998-06-04
PRIOR APPLICATION NUMBER: 60/088167 PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR FILING DATE: 1998-06
APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: 60/088029 FILING DATE: 1998-06-04 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/087759 FILING DATE: APPLICATION NUMBER: 60/087609 FILING DATE: 1998-05-28 FILING DATE: 1998-01
APPLICATION NUMBER: APPLICATION NUMBER: 60/088861 FILING DATE: 1998-06-11 APPLICATION NUMBER: 60/088876 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088738 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088655 FILING DATE: 1998-06-09 APPLICATION NUMBER: 60/088734 FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088217 FILING DATE: 1998-06-05 FILING DATE: 1998-06-05
APPLICATION NUMBER: 60/088202
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FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088326 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088030 APPLICATION NUMBER: 60/088026 FILING DATE: APPLICATION NUMBER: 60/088025 APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/087827 FILING DATE: APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: 1998-06-11 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-12 APPLICATION NUMBER: 60/088824 APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: APPLICATION NUMBER: 60/088810 APPLICATION NUMBER: 60 FILING DATE: 1998-06-1 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089514 FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089512 APPLICATION NUMBER: 60/089440 FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089599 APPLICATION NUMBER: 60/089532 FILING DATE: FILING DATE: 1998-06-04 1998-06-04 1998-06-02 1998-06-02 1998-06-04 1998-06-03 1998-06-02 1998-06-11 1998-06-10 1998-06-10 1998-06-04 1998-06-10 1998-06-16 1998-06-17 60/088033 60/088028 60/088021 60/087106 60/087607 60/088861 60/088858 60/088742 60/089598 6-17 60/089105 60/088826 60/089538 60/089600

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FILING DATE: 1998-06-26
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APPLICATION NUMBER: 60/091633
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APPLICATION NUMBER: 60/090676
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FILING DATE: 1998-06-24
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FILING DATE: 1998-06-24
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FILING DATE: 1998-06-23
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                                                                                   PRIOR FILING DATE: 1998-0
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1998-0
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PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Secreted and Transmembrane Polypeptides TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730PIC19
CURRENT APPLICATION NUMBER: US/09/990,444
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR PILING DATE: 1997-06-16
PRIOR PILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
DBIOD APPLICATION NUMBER: 60/062250
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PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
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APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
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PRIOR ETLING DATE: 1998-07-07
PRIOR APPLICATION NUMBER: 60/091982
PRIOR FILING DATE: 1998-07-07
PRIOR APPLICATION NUMBER: 60/092182
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FILING DATE: APPLICATION N
                             APPLICATION NUMBER: 60/087607
                                                APPLICATION NUMBER: 60/
FILING DATE: 1998-05-28
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Kljavin, Ivar J.
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D. US20020193300A1
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                1998-06-02
                                                                                     1998-05-07
                                                                                                                   1998-04-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Timothy A.
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Pred. No. 1.5e+05;
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PRIOR DR APPLICATION NUMBER: 60/088861
DR FILING DATE: 1998-06-11
DR APPLICATION NUMBER: 60/08876
DR FILING DATE: 1998-06-11
DR APPLICATION NUMBER: 60/089105
DR FILING DATE: 1998-06-12
DR FILING DATE: 1998-06-12 DR FILING DATE: 1998-06-17

PR APPLICATION NUMBER: 60/089653

PR FILING DATE: 1998-06-17

PR APPLICATION NUMBER: 60/089801

PR FILING DATE: 1998-06-18

PR APPLICATION NUMBER: 60/089907 FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088026
FILING DATE: 1998-06-04 FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/087759 FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/088033 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088326 FILING DATE: 1998-06-04 FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088030
FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088028 APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/088858 FILING DATE: 1998-06-11 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088826 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-10 FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088655 FILING DATE: 1998-06-09 APPLICATION NUMBER: FILING DATE: 1998-06 APPLICATION NUMBER: 60/088167 APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: FILING DATE: 1998-06-03 APPLICATION NUMBER: FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/ APPLICATION NUMBER: 60/ FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/(FILING DATE: 1998-06-10 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088202 FILING DATE: APPLICATION NUMBER: 60/089598 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089538 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/088824 FILING DATE: APPLICATION NUMBER: 60/089600 FILING DATE: APPLICATION NUMBER: FILING DATE: 1998-06-17 FILING DATE: APPLICATION NUMBER: 60/089512 1998-06-04 1998-06-04 1998-06-10 1998-06-10 1998-06-1 1998-06-05 1998-06-05 1998-06-17 60/088029 60/088025 60/088021 60/087827 60/089514 60/089440 60/088810 60/088742 60/088738 60/088734 60/088217 60/088212 60/089599

PRIOR DR APPLICATION NUMBER: 60/090863
R FILING DATE: 1998-06-26
DR APPLICATION NUMBER: 60/091360
R FILING DATE: 1998-07-01
DR APPLICATION NUMBER: 60/091478
DR FILING DATE: 1998-07-02 APPLICATION NUMBER: 60/089908 FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/089947 FILING DATE: 1998-06-24
APPLICATION NUMBER: 60/090535
FILING DATE: 1998-06-24
APPLICATION NUMBER: 60/090540
FILING DATE: 1998-06-24 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-24 FILING DATE: 1998-06-22 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-22 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-22 APPLICATION NUMBER: 6 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 FILING DATE: APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: 60/089952 FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/090542 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/090557 FILING DATE: 1998-06-24 APPLICATION NUMBER: APPLICATION NUMBER: 60/090355 FILING DATE: FILING DATE: APPLICATION NUMBER: 60/090862 FILING DATE: 1998-06-26 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-25 FILING DATE: 1998-06-25 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-25 APPLICATION NUMBER: 60/090678 APPLICATION NUMBER: APPLICATION NUMBER: 60/090349 APPLICATION NUMBER: 60/090246 FILING DATE: 1998-07-07 APPLICATION NUMBER: 60/092182 APPLICATION NUMBER: 60/091633 FILING DATE: 1998-07-02 APPLICATION NUMBER: 60/091519 FILING DATE: 1998-07-02 APPLICATION NUMBER: 60/091544 FILING DATE: 1998-07-01 FILING DATE: 1998-00 APPLICATION NUMBER: APPLICATION NUMBER: 60/090695 APPLICATION NUMBER: 60/090676 FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: 60/091978 FILING DATE: 1998-07-02 APPLICATION NUMBER: 60/091626 FILING DATE: 1998-06-25 1998-06-24 1998-06-23 1998-06-1998-06-1998-06-19 1998-06-24 1998-07-07 1998-06-25 -90-8661 60/090472 60/090445 60/090435 60/090431 60/090254 60/090252 60/089948 60/091982 60/090696 60/090694 069060709 60/090444 60/090429

Query Match

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                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
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                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/066770
PRIOR FILING DATE: 1997-11-24
                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P273091C69
CURRENT APPLICATION NUMBER: US/09/989,730
CURRENT FILING DATE: 2001-11-20
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APPLICANT:
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                          APPLICATION NUMBER: 60/
FILING DATE: 1998-06-03
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Stewart, Timothy A.
Tumas, Daniel
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Williams,
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Kljavin, Ivar J.
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Botstein, David
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PRIOR APPLICATION NUMBER: 60/088824 APPLICATION NUMBER: 60/089947 FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/089599 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089948 APPLICATION NUMBER: 60/089532 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/089105 FILING DATE: 1998-06-12 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-11 APPLICATION NUMBER: 60/088810 FILING DATE: 1998-06-10 FILING DATE: 1998-06-09 APPLICATION NUMBER: 60/088734 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088738 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088217 APPLICATION NUMBER: 60/088212 FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088202 FILING DATE: 1998-06-05 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088033 FILING DATE: 1998-06-04 1998-06-18 1998-06-18 1998-06-19 1998-06-18 1998-06-17 1998-06-17 1998-06-17 1998-06-1 1998-06-16 1998-06-16 1998-06-1 1998-06-10 1998-06-16 1998-06-1 1998-06-10 1998-06-10 1998-06-10 1998-06-10 1998-06-04 1998-06-05 1998-06-05 1998-06-05 1998-06-04 1998-06-1998-06-04 60/089952 60/089907 60/089801 806680/09 60/089653 60/089600 60/089598 60/089538 60/089514 60/089512 60/089440 60/088876 60/088861 60/088858 60/088826 60/088742 60/088655 60/088167 60/088326 60/088030 60/088029 60/088028 60/088026 60/088025

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FILING DATE: 1998-06-24
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APPLICATION NUMBER: 60/088021
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APPLICATION NUMBER: 60/087607
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Grimaldi, J. Christopher
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Stewart, Timothy A.
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APPLICATION NUMBER: 60/088861
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APPLICATION NUMBER: 60/089512
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APPLICATION NUMBER: 60/090694
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Sequence 122, Application US/09991181 Publication No. US20020197615A1 GENERAL INFORMATION:

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Gaps

Ashkenazi, Avi J. Baker, Kevin P.

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CURRENT APPLICATION NUMBER: US/09/991,181
CURRENT FILING DATE: 2001-11-16
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OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/065186
OR FILING DATE: 1997-11-12
OR APPLICATION NUMBER: 60/065311
OR FILING DATE: 1997-11-13
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FILING DATE: 1998-05-28
APPLICATION NUMBER: 60/1
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APPLICATION NUMBER: 60/088326
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FILING DATE: 1997-11-24
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APPLICATION NUMBER: 60/088021
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APPLICATION NUMBER: 60/088025
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APPLICATION NUMBER: 60/087759
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APPLICATION NUMBER: 60/088033
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Godowski, Paul J.
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PRIOR APPLICATION NUMBER: 60/088167 PRIOR FILING DATE: 1998-06-05 PRIOR APPLICATION NUMBER: 60/088202 PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: 60/088738
PRIOR FILING DATE: 1998-06-10 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-0 PRIOR FILING DATE:
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DR FILING DATE: 1998-06-16
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APPLICATION NUMBER: 60/08876
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APPLICATION NUMBER: 60/089105
FILING DATE: 1998-06-12 APPLICATION NUMBER: 60/088824 FILING DATE: 1998-06-10 APPLICATION NUMBER: FILING DATE: 1998-06 APPLICATION NUMBER: 60/088742 FILING DATE: 1998-06-10 FILING DATE: 1998-06-09 APPLICATION NUMBER: 60/088655 FILING DATE: 1998-06-11 APPLICATION NUMBER: 60/ APPLICATION NUMBER: 60/088858 APPLICATION NUMBER: 60/088826 FILING DATE: 1998-06-10 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/089440 FILING DATE: 1998-06-16 FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/089952 FILING DATE: 1998-06-19 APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: 60/089598 FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/090355 FILING DATE: 1998-06-23 APPLICATION NUMBER: 60/090349 FILING DATE: 1998-06-23 APPLICATION NUMBER: APPLICATION NUMBER: 60/089948 FILING DATE: 1998-06-19 FILING DATE: APPLICATION NUMBER: 60/089947 APPLICATION NUMBER: 60/089908 FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/089653 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-06-22 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-22 FILING DATE: 1998-06-22 APPLICATION NUMBER: 1998-06-05 1998-06-17 1998-06-10 1998-06-05 1998-06-17 1998-06-10 1998-06-24 1998-06-19 60/088734 60/088212 60/088810 60/089801 60/089600 60/089599 60/089538 60/089532 5-17 60/089514 60/089512 60/088861 60/089907 60/090429 60/090254 60/090252 60/090246

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RESULT 20
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                                           Sequence 122, Application US/09993687
Publication No. US20020198149A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J
APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
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                                  APPLICANT:
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PRIOR APPLICATION NUMBER: 60/091978
PRIOR FILING DATE: 1998-07-07
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PRIOR FILING DATE: 1998-06-26
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APPLICATION NUMBER: 60/092182
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APPLICATION NUMBER: 60/091519
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FILING DATE: 1998-06-25
APPLICATION NUMBER: 60/090690
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APPLICATION NUMBER: 60/090694
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Fong, Sherman
                Ferrara, Napoleone
                                  Eaton, Dan L.
                                                                                                                                                                                                                                                                                                          Conservative
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PRIOR APPLICATION NUMBER: 60/065311
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PRIOR FILING DATE: 1998-04-28
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CURRENT FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/049787
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PRIOR APPLICATION NUMBER: 60/062250
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C11
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APPLICATION NUMBER: 60/088217
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FILING DATE: 1998-06-04
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Stewart, Timothy A.
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Gerritsen, Mary E.
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DR FILING DATE: 1998-06-17
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FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088824
FILING DATE: 1998-06-10
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APPLICATION NUMBER: 60/089512
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APPLICATION NUMBER: 60/090431
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APPLICATION NUMBER: 60/090444 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/090445
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RESULT 21
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DR FILING DATE: 1998-06-25
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FILING DATE: 1998-06-24
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Fong, Sherman
Gerber, Hanspeter
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Kljavin, Ivar J.
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                                                                                                                                                                                   Botstein, David
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TITLE OF INVENTION: Acids Encoding the Si
FILE REFERENCE: P2730P1C64
CURRENT APPLICATION NUMBER: US/09/989,734
CURRENT FILING DATE: 2001-11-19
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FILING DATE: 1998-06-09
APPLICATION NUMBER: 60/088734
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APPLICATION NUMBER: 60/088738
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APPLICATION NUMBER: 60/088217
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APPLICATION NUMBER: 60/066770
FILING DATE: 1997-11-24
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FILING DATE: 1997-06-16
APPLICATION NUMBER: 60/062250
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PR APPLICATION NUMBER: 60/091478
DR FILING DATE: 1998-07-02
DR APPLICATION NUMBER: 60/091544
DR FILING DATE: 1998-07-01
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FILING DATE: 1998-06-25
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APPLICATION NUMBER: 60/091982
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FILING DATE: 1998-07-02
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FILING DATE: 1998-07-02
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29; Conservative
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   Roy,
Stewart, 1...
Stewart, 1...
Col
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Godowski, Paul J.
                                                                                                                                                                                              Fong, Sherman
Gerber, Hanspeter
                                                                                                                                                                                                                                                                       Botstein, David
                                                                                                                                                                                                                                                                                          Baker, Kevin P
                         Paoni, Nicholas F.
Roy, Margaret Ann
Stewart, Timothy A.
                                                                                                                                                                                 Gerritsen, Mary E.
                                                                                                                                                                                                                              Ferrara, Napoleone
                                                                                                                                                                                                                                              Eaton, Dan L.
                                                                                                                                                                                                                                                           Desnoyers, Luc
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                                                                                                       Kljavin,Ivar J.
                                                                                                                    Gurney, Austin_L.
                                                                                                                                   Grimaldi, J. Christopher
                                                                            Pan, James
                                                                                       Napier, Mary A.
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   Colin K.
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Pred. No. 1.5e+05;
O; Mismatches 15; Indels
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PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-12
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CURRENT FILING DATE: 2001-11-15
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PRIOR APPLICATION NUMBER: 60/088026
PRIOR FILING DATE: 1998-06-04
PRIOR APPLICATION NUMBER: 60/088028
PRIOR FILING DATE: 1998-06-04
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-02-25
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PRIOR FILING DATE: 1998-06-02
PRIOR APPLICATION NUMBER: 60/087759
PRIOR FILING DATE: 1998-06-02
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PRIOR APPLICATION NUMBER: 60/087607
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APPLICATION NUMBER: 60/
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APPLICATION NUMBER: 60/088030
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APPLICATION NUMBER: 60/0
FILING DATE: 1998-06-10
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FILING DATE: 1998-06-23
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APPLICATION NUMBER: 60/089653
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APPLICATION NUMBER: 60/089598
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Best Local
         TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730P1C4
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FILING DATE: 1998-07-02
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APPLICATION NUMBER: 60/091633
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APPLICATION NUMBER: US/09/993,667
                                                         Zhang, Zemin
                                                                     Watanabe, Colin K.
Williams, P. Mickey
Wood, William I.
                                                                                                                          Roy, Margaret Ann
Stewart, Timothy A.
                                                                                                                  Tumas,
                                                                                                                                                          Paoni, Nicholas F.
                                                                                                                                                                                                   Kljavin,Ivar
                                                                                                                                                                                                                  Gurney, Austin L.
                                                                                                                                                                                                                              Grimaldi, J. Christopher
                                                                                                                                                                                                                                           Goddard, Audrey
Godowski, Paul
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                                                                                                                                                                          Pan,James
                                                                                                                                                                                   Napier, Mary A.
                                                                                                                                                                                                                                                                                                                                  Eaton, Dan L
                                                                                                                                                                                                                                                                                                                                                           Botstein, David
                                                                                                                                                                                                                                                                                                                                                                          Baker, Kevin P.
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Pred. No.
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CURRENT FILING DATE: 2001-11-14 PRIOR APPLICATION NUMBER: 60/049787 OR APPLICATION NUMBER: 60/088026
OR FILING DATE: 1998-06-04
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R FILING DATE: 1998-06-10
R APPLICATION NUMBER: 60/088858
R FILING DATE: 1998-06-11
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R FILING DATE: 1998-06-11
R FILING DATE: 1998-06-11 FILING DATE: 1997-06-16 APPLICATION NUMBER: 60/ FILING DATE: 1997-10-17 APPLICATION NUMBER: 60/087609 FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/087759 FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/084600 FILING DATE: 1998-05-07 APPLICATION NUMBER: 60/087106 APPLICATION NUMBER: 60/075945 FILING DATE: 1998-02-25 APPLICATION NUMBER: 6 FILING DATE: 1997-11-APPLICATION NUMBER: 60/065186 FILING DATE: 1997-11-12 APPLICATION NUMBER: 60/088326 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088167 FILING DATE: 1998-06-05 APPLICATION NUMBER: 60/088202 FILING DATE: 1998-05-28
APPLICATION NUMBER: 60/087607
FILING DATE: 1998-06-02 APPLICATION NUMBER: 60/083322 FILING DATE: 1998-00 APPLICATION NUMBER: FILING DATE: 1998-06-09 APPLICATION NUMBER: 60/088734 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088033 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088021 APPLICATION NUMBER: 60/087827 FILING DATE: 1998-06-03 FILING DATE: 1998-04-28 APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: 60/088742 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/088212 FILING DATE: 1998-06-05 FILING DATE: 1998-06-04 FILING DATE: FILING FILING DATE: APPLICATION NUMBER: 60/088810 FILING DATE: 1998-06-10 APPLICATION NUMBER: APPLICATION NUMBER: 60/088217 FILING DATE: 1998-06-05 APPLICATION NUMBER: APPLICATION NUMBER: 60/088655 FILING DATE: 1998-05-28 1998-03-20 1997-11-13 1998-06-10 1998-06-05 1998-06-10 1998-06-04 2001-11-14 60/078910 60/066770 60/065311 60/088738 60/088326 60/088025 60/088824 /062250

PRIOR R FILING DATE: 1998-06R APPLICATION NUMBER: (
R FILING DATE: 1998-06R FILING DATE: 1998-06R FILING DATE: 1998-06-APPLICATION NUMBER: 60/089105 FILING DATE: 1998-06-12 APPLICATION NUMBER: 60/089440 FILING DATE: 1998-06-16 FILING DATE: 1998-06-19
APPLICATION NUMBER: 60/090246
FILING DATE: 1998-06-22 FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-18 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-17 FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/089532 FILING DATE: 1998-06-23
APPLICATION NUMBER: 60/090429
FILING DATE: 1998-06-24
APPLICATION NUMBER: 60/090431
FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/089947 FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/ APPLICATION NUMBER: 60/ FILING DATE: 1998-06-17 APPLICATION NUMBER: FILING DATE: 1998-00 APPLICATION NUMBER: 60/089598 FILING DATE: FILING DATE: APPLICATION NUMBER: 60/089514 FILING DATE: FILING DATE: 1998-06-22
APPLICATION NUMBER: 60/090349
FILING DATE: 1998-06-23 FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: 60/090695 FILING DATE: 1998-06-25 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/ APPLICATION NUMBER: APPLICATION NUMBER: 60/090535 FILING DATE: 1998-06-24 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/090472 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/090435 APPLICATION NUMBER: 60/090355 APPLICATION NUMBER: APPLICATION NUMBER: 60/090252 APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-06-24 FILING DATE: 1998-06-24 APPLICATION NUMBER: 60/090445 FILING DATE: FILING DATE: 1998-06-25 NUMBER: 60/089600 1998-06-17 1998-06-1998-06-1998-06-24 1998-06-22 1998-06-19 1998-06-1998-06-25 1998-06-25 1998-06-25 1998-06-24 60/089538)6-17 60/090254 60/089948 60/089907 60/089801 60/089653 60/089599 60/089512 60/090444 60/089952 60/090690 60/090678 60/090676 60/090557 60/090542 60/090540 806680/09 60/090696 60/090694

FILING DATE: 1998-06-25

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US-09-990-438-122

Sequence 122, Application US/09990438

Publication No. US20030027754A1

GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: US/09/561,490
PRIOR FILING DATE: 2000-04-28
PRIOR APPLICATION NUMBER: US 60/131,448
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: LOCALIZATION OF MAJOR PEPTIDE AUTOEPITOPES FOR NUCLEOSOME SPECIF TITLE OF INVENTION: IC T CELLS OF SYSTEMIC LUPUS ERYTHEMATOSUS FILE REFERENCE: 290-13U1 5662-5017
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APPLICANT: KALIYAPERUMAL, Arunan
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PRIOR FILING DATE: 1998-07-07
PRIOR APPLICATION NUMBER: 60/091982
PRIOR FILING DATE: 1998-07-07
PRIOR APPLICATION NUMBER: 60/092182
                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Histone fragment
                                                                                                                                            1260 ACAGCTGGGCGCACCGGGCCTTCGGCCCTTCGGCCGAC 1303
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FILING DATE: 1998-07-02
APPLICATION NUMBER: 60/091626
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FILING DATE: 1998-07-02
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APPLICATION NUMBER: 60/091478
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Pred. No. 1.5e+05;
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PRIOR APPLICATION NUMBER: 60/066770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
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PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
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PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
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CURRENT APPLICATION NUMBER: US/09/990,438

CURRENT FILING DATE: 2001-11-14
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APPLICATION NUMBER: 60/
FILING DATE: 1998-06-04
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Williams, P. Mickey
Wood, William I.
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Stewart, Timothy A.
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APPLICATION NUMBER: 60/088167
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FILING DATE: 1998-06-17
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APPLICATION NUMBER: 60/091360
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Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki

Toshiyuki and Piero Carninci (RIKEN)

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                         1 (bases 1 to 43)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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BI462985.1 GI:15253641
                                                                                                                                                                                                                     BI462985 43 bp mRNA linear EST 21-AUG-2603204504F1 NIH_MGC_97 Homo sapiens cDNA clone IMAGE:5270424 5',
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Best Local Similarity
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 50)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly

M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.,

and Wright, D., Weiss, R.
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Contact: Robert B.
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                                                                                                                                                            High quality sequence stop: 50.
Location/Qualifiers
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Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Errc
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Fax: 801 585 7177
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Tissue Procurement: Life Technologies,
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BI772215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 bp mRNA linear EST 25-SEP-2001
603056165F1 NIH_MGC_122 Homo sapiens cDNA clone IMAGE:5205425 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nttp://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BI772215.1 GI:15763793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA Library Preparation: Life Technologies, Inc. (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (bases 1 to 45)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNRD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                           /organism-"Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5205425"
/clone_lib="NIH_MGC_122"
/lab_host="DH10B"
/note="Organ: pooled lung and spleen; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: EcoRV (destroyed); RNA source anonymous pool of 24 week female lung, 16 week female spleen, and 20-22 week male spleens. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.4 kb, insert size
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.6%;
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Pred. No. 2.5e+05;
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VERSION KEYWORDS

ACCESSION DEFINITION

SOURCE

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COMMENT

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TITLE AUTHORS

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ORIGIN

Matches

BASE COUNT

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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Blaxter ML Institute of Cell, Animal and Population Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genes expressed in adult female Brugia malayi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blaxter, M.L., Waterfall, M., Daub, J., Lizotte, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brugia malayi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MBAFCEZDO3T3 Brugia malayi adult female cDNA (SAW96MLW-BmAF) Brugia malayi cDNA clone AFCFZDO3 5', mRNA sequence.
                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The ABI trace of this sequence can be viewed at http://www.sanger.ac.uk/brugia/AFC/MBAFCF2D03T3.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: mark.blaxter@ed.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: +44 131 650 6760
Fax: +44 131 670 5450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ashworth Labs, King's Buildings, West Mains Road, Edinburgh, EH9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Edinburgh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Onchocercidae; Brugia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Nematoda; Chromadorea; Spirurida; Filarioidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brugia malayi.
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а
                                                                                            library is available from genome@smith.edu."
a 23 c 2 g 24
                                                                                                                                                                                                                                       /lab_host="XL1-Blue MRF/"
/note="Vector: Lambda Uni-ZAP XR; Site_1: Eco RI; Site_2:
Xho I; Lymphatic filarial nematode parasite of humans.
MRNA was prepared from approximately 50 adult females
isolated from the peritoneal cavity of jirds and
                                                                                                                                                  transcriptase and oligo(dT) followed by RNase H and DNA pol I. The library has 5 x 10E6 independent recombinants and the average insert size is -900bp.The library was constructed by Michelle Lizotte-Waniewski. The
                                                                                                                                                                                                                               converted to double-stranded cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 026. this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                             /dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                   /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="AFCF2D03"
                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:6279"
                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Brugia malayi"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                   Clone_11b="Brugia malayi adult female cDNA (SAW96MLW-BmAF
                       1.6%;
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                       Pred. No. 3.8e+05;
                                       Score 25;
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Pred. No.
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                                                                                                  24 t
                                           DB 9;
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                                       Length 49;
                                                                                                                                    Dr.S.A.Williams, email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 45
   Indels
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DEFINITION
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AI813244
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AV833587/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 GAGAGCGAGCGGGCGAGCCGAGCGAAGGGAAAGCGCAAGAGAGAG 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              34;
                  Unpublished (2001)
Contact: Kazuhiro Sato
Research Institute for Bioresources
                                                                                                                                                Hordeum vulgare subsp. vulgare.
Hordeum vulgare subsp. vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooidea
                                                                                                                                                                                                                                                                  vulgare shoots germination Hordeum vulgare clone bags16a08, mRNA sequence.

AV833587
                                                                                                                                                                                                                                                                                                                            AV833587 K. Sato unpublished cDNA library:
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1 (bases 1 to 49)
Whetten, R.W., Kinlaw, C.S., Retzel, E. and Sederoff, R.R. The Pine Gene Discovery Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pinus taeda
                                                                                                                                                                                                                                  EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: rosswhet@unity.ncsu.edu
Seq primer: T3.
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EST.
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Okayama University, Barley Germplasm Center
                                                                        Barley EST sequencing project in NIG and Okayama Univ
                                                                                                  Sato,
                                                                                                                                                                                                                                                    AV833587.1 GI:14525676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fax: 919-515-7801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Forest Biotechnology Group
North Carolina State University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Ross Whetten
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Coniferopsida; Coniferales; Pinaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              loblolly pine.
                                                                                                                                    Triticeae;
                                                                                                                  (bases 1 to 49)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . of Forestry, NC State University, 6113 Jordan Hall, Raleigh 27695-8008 919-515-7800
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Pine Lambda Zap Xylem library"
/tissue_type="differentiating xylem"
/note="Vector: Lambda Zap; Site_1: EcoRT; Site_2: XhoI;
/note="Vector: Lambda Zap; Site_1: EcoRT; Site_2: XhoI;
Differentiating xylem was collected from the main stem a 35-year old lobloly pine tree harvested during the growing season. RNA isolation and library preparation followed the methods of Allona et al., PNAS 95:9693-8,
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                                                                                                                                      Hordeum.
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Pinus; Pinus.
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Best Local Similarity
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Sato,K., Saisho,D., Takeda,K., Shini,T. and Kohara,Y. Direct
                                                                                                                                                                                                                                                                                                                                                                                      found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM10005 row: i column: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Email: kazsato@rib.okayama-u.ac.jp,
                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
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                                                                            /tissue_type="adrenal cortex carcinoma, cell line" /lab_host="DH10B (phage-resistant)" /note="organ: adrenal gland; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: SalI; Cloned unidirectionally; oligo-dT primed. Average insert size 1.29 kb. Library enriched for full-length clones and constructed by Life Technologies. Note: this is a NHH_MGC Library."
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vulgare subsp. vulgare shoots germination"
/tissue_type="shoots"
/tissue_type="shoots"
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/cultivar="Haruna Nijo"
/db_xref="taxon:112509"
                                                                                                                                                                                                                                           /clone="IMAGE:4361943"
/clone_lib="NIH_MGC_84"
                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
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25 c 0 q 2
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Kohara,Y., Shin-i,T., Kimura,T., Nedara, Kohara,Y., Shin-i,T., Kimura,T., Nedara, Kanara, Kana
                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Atherinomorpha; Beloniformes; Adrianichthyidae; Oryzinae; Oryzias.
                                         Center For Genetic Resource Information
                                                                                                                                                                                                                                                                                         Oryzias latipes
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                                                                     Contact: Tadasu Shin-i
                                                                                                  Unpublished (2001)
                                                                                                                                                                                                                                                                                                                   Japanese medaka.
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                       National Institute of Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                         BJ000572 MF01SSA cDNA Oryzias
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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AV947640 Nori Satoh unpublished cDNA library, young adult Ciona
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Department of Zoology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: satoh@ascidian.zool.kyoto-u.ac.jp
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/db_xref="taxon:7719"
/clone="ciad01g16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="whole animal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Nori Satoh unpublished cDNA library, young
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Pred. No. 5.4e+05;
D; Mismatches 14
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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Location/Qualifiers
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/clone_lib="NCI_CGAP_Panl"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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/db_xref="taxon:8090"
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DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
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NGI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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Tel: 81-559-81-6856
Fax: 81-559-81-6855
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dr. Average insert size 1.72 kb. Life Technologies catalog #: 11548-013"
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/tissue_type="whole embryo"
/dev_stage="segmentation stage 20 - 25"
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Department of Zoology
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                                                     Satoh,N., Satou,Y., Kohara,Y. and Shin-i,T.
Expressed genes in Ciona intestinalis
Unpublished (2000)
Contact: Nori Satoh
Department of Zoology
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             Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
                                                                                                                             Phlebobranchia; Cionidae; Ciona.
1 (bases 1 to 47)
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Fax: 81-75-705-1113
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/dev_stage="egg"
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/tissue_type="whole_animal"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ385990.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eutheria;
                                                                                                                                                    /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GI:10499690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.5%;
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12 AGCGAGGGAGAGCGAGCGGGGGAGCGAGGGAAGGGAAAG
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Insert Length: 10000 Std Error: (
Plate: 0144 row: D column: 16
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 47)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          clone UUGC1M0144D16 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ385990 47 bp DNA linear, GSS 02-OCT-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     308, Biomedical Polymers Research Bldg., 20 S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: pBluescript SK"
23 c 0 g 23 t
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/tissue_type="whole animal"
/dev_stage="larva"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:7719"
/clone="cilv02g12"
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                   was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                  /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                /db_xref="taxon:10090"
/clone="UUGC1M0144D16"
                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                              (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                    Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                    /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                    /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus'
ligated to the blunt ends in high
                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 23.6; DB 10;
Pred. No. 7e+05;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
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     molar excess.
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                                                                                                                                         The DNA
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RESULT 14
AZ853064/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0156 row: K column: 04
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ853064
2M0156K04F Mouse 10kb plasmid UUGCCIM library Mus musculus clone UUGC2M0156K04 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 47)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                 /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack:
                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC2M0156K04"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                               /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                              /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                 /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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REFERENCE

AUTHORS

KEYWORDS VERSION ACCESSION DEFINITION FOCUS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0170 row: N column: 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: 0170 row: N column: 09 Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
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801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114[gb]AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
/clone="UUGC2M0170N09"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampisillin resistance. The color of the co
                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                       /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             69.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0174 row: M column: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                     /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:10090"
/clone="UUGC2M0174M10"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                         /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                            /sex="Male"
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Taylor,R., Ashurst,J.L., Croning,M.D.R., Zu
Sanger Xenopus tropicalis EST project 2002
Unpublished (2001)
   30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Taylor R
Sanger Centre
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Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Silurana tropicalis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          constructed by Aaron M. Zorn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequencing primer: PICSP6
This sequence is from a Xenopus Gene Collection (XGC) library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      western clawed frog.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: trop@sanger.ac.uk
Sanger Xenopus tropicalis EST project 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Xenopodinae; Silurana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hinxton, Cambridgeshire, CB10 1SA, UK
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                                                                                                                                                                                                                        /dev_stage="gastrula (stages 10.5-13 mixed)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="XGC-gastrula"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="TGas053o17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Silurana tropicalis"
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                                    Score 23.4; DB 9; Pred. No. 7.6e+05;
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                                                                             DB 9;
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ORGANISM
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                        source
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High quality sequence stop: 49
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84112,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
Contact: Robert B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 49)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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                                                                                              Conservative
                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 jgb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Mouse 10kb plasmid UUGC1M library"
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81.8%;
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Pred. No. 7.7e+05;
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                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                     AV672475 Nori Satoh unpublished cDNA library Ciona intestinalis cDNA clone citb5m21 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Library was constructed by Debopam Chakrabarti DNA sequencing by: Washington University Genome Sequencing Center For information on obtaining a clone please contact: L. David Sibley (sibley@borcim.wmstl.edu), Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tang, K., Cole, R., Chakrabarti, D., Haywood, R., Clifton, S., Pape, D. Marra, M., Hillier, L., Martin, J., Wylie, T., Dante, M., Theising, B. Bowers, Y., Gibbons, M., Ritter, E., Bennett, J., Jentes, E., Ronko, I. Tsagareishvili, R., Belaygorod, L., Franklin, C., Carr, L., Grow, A., Maguire, L., Richey, J., Wadkins, J., Kennedy, S., Levinso, D., Waterston, R., Wilson, R. and Sibley, D. Waterston, R., Wilson, R. and Sibley, D. WashU Plasmodium EST Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
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Fax: 314 286 1810
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Contact: L. David Sibley
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                                                                                                                                                                                                                                                                                                                                                                                                                           /Clones were mass excised using the Exassist helper phage (5to and extracted with phenol/chlores were mass excised using the Exast the Econa of the Cona in the phagemid was electroporated into Dhiob end extracted with phenol/chloroform method. The poly A+ RNA was isolated by the polyAT-Tract mana isolation system (promega, WI) using streptavidin Magnesphere particles. Directional CDNA Hibraries were constructed by oligo d(T) priming of poly(A)+ RNA (5mg) into EconI and XhoI sites of 1 ZapII vector using the Zap CDNA inserts in the library was between 1.0 and 1.5kb. Clones were mass excised using the Exassist helper phage (Stratagene), the phagemids were precliptated with pEG 8000 and extracted with phenol/chloroform. Phagemid DNA was electroporated into DH10B cells."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:5833"
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                                                                                                                                                                                                                                                                                                                                         1.5%;
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                                                                                                                                                                                                                                                                                                                                   Score 23.4; DB 14; Pred. No. 7.7e+05;
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                                                                       EST 05-OCT-2000
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                                                                                                                                                                                                                                                                                                           Gaps
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В QΥ

VERSION

AV672475.1 GI:10110474

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REFERENCE
AUTHORS
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AA977986/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA977986 49 bp mRNA linear EST 26-MAY-1998 oq55a06.sl NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:1590226 3' similar to TR:000301 000301 KSRP. ;contains element MER22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    repetitive element; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Nori Satoh
Department of Zoology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T. Expressed genes in Ciona intestinalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phlebobranchia; Cionidae; Ciona.
1 (bases 1 to 44)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 49)
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                                                              High quality sequence stop: 1.
Location/Qualifiers
                                                                                                   Trace considered overall poor quality Seq primer: -40ml3 fwd. ET from Amers
                                                                                                                                                                 cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                        Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA977986.1 GI:3155432
                                                                                                                                                                                                                                                                                                  Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
                                                                                                                                                                                                                                                                            cDNA Library Preparation: M. Bento Soares, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81-75-705-1113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Nori Satoh unpublished cDNA library"
/tissue_type="whole animal"
/dev_stage="tailbud"
/note="Vector: pBluescript SK"
22 c 1 g 21 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Ciona intestinalis"
/db_xref="taxon:7719"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 8.3e+05;
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1343 GGCGGGGACAGCGGCGGCGGGGACCGCGG 1371

GGCGGCGACAGCGGCGGCTGGGCTCGCGG

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BASE COUNT
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
                                                                                                                                                                                                                                                                                                                                Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedal, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HEP05140, mRNA sequence. AU103717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AU103717 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                         ,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                             Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia;
                    Similarity
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Conservative
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                                                                                               dimethylfumarate treated 20 c 16 g
                                                                                                                  /clone_lib="Sugano Homo sapiens cDNA library" /note="Differential display comparison of untreated and dimethylfumarate treated_U937 cells"
                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                             /clone="HEP05140"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                    ocation/Qualifiers
                  1.4%;
86.2%;
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  0; Mismatches
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Pred. No. 1.1e+06;
                  Score 22.6; DB 9;
Pred. No. 1.1e+06;
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                                          DB 9;
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                                                     22 AGAGCGAGCGGGGGGAGCGAGGGAAGGGGAAAGCGCAAGAGAGAG 69
                                                                                                                                        Loca l
  w
AGAGAGGGAAGAAGAGGAAGGGAAGGGGAGAGAGGGAAAG 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          National Institute on Aging/National Institutes of Health 333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820, USA Email: cdnaelgsun.grc.nia.nih.gov
This clone set has been freely distributed to the community. Please visit http://lgsun.grc.nia.nih.gov/cDNA/15k.html for details. Plate: H3022 row: G column: 12
Seq primer: -21M13 Reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: George J. Kargul Laboratory of Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2001)
Other_ESTs: H3022G12-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kargul, G.J., Dudekula, D.B., Qian, Y., Lim, M.K., Jaradat, S.A., Tanaka, T.S., Carter, M.G. and Ko, M.S.H.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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BG078033
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                                                                                                                                                                                                                                                         18
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                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="vector: psport1; Site_1: Sal1; Site_2: Not1; This /note="vector: psport1; Site_1: Sal1; Site_2: Not1; This clone is among a rearrayed set of 15,247 clones from 11 embryo cDNA libraries (including preimplantation stage embryo cDNA libraries (including preimplantation stage embryos from unfertilized egg to blastocyst, embryonic part of E7.5 embryos, extraembryonic part of E7.5 embryos, and cDNA library. Average insert size 1.5 kb. All source libraries are cloned unidirectionally with Oligo(dT) 'Not primers. References include: (1) Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray, 2000, Proc. Natl. Acad. Sci. U S A, 97: 9127-9132; (2) Large-scale cDNA analysis reveals phased gene expression patterns during preimplantation mouses developments.
                                                                                                                                                                                                                                                                                                                                   patterns during preimplantation mouse development, 2000, bevelopment, 127: 1737-1749; (3) Genome-wide mapping of unselected transcripts from extraembryonic tissue of 7.5-day mouse embryos reveals enrichment in the t-complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="NIA Mouse 15K cDNA Clone Set"
/sex="Clones arrayed from a variety of cDNA libraries"
/dev_stage="Clones arrayed from a variety of cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /strain="C57BL/6J"
/db_xref="niaEST:H3022G12-5"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                   under-representation on the X chromosome, 1998, Hum Genet 7: 1967-1978." 0 c 31 g 1 t
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                                                                                                                                        1.48;
                                                                                                                                        Score 22.4;
Pred. No. 1.
                                                                                                             Mismatches
                                                                                                                                        .2e+06;
                                                                                                                                                                   DB 12;
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RESULT 25
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                                                                                                             JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
Toshiyuki and Piero Carninci (RIKEN)
                                                                                                        NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                        B1547391 43 bp mRNA linear EST 05-SEP-2001 603197282F1 NIH_MGC_95 Homo sapiens cDNA clone IMAGE:5276516 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA Sequencing by: National Institutes of Health Intramural Sequencing Center (NISC)
Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: info@image.llnl.gov
Plate: LLAM11840 row: N column: 1
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                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 43)
                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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1 (bases 1 to 50)
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/lab_host="DH10B (phage-resistant)"
/note="Organ: brain; vector: pCMV-SPORT6.ccdb; Site_1
/note="Organ: brain; vector: pCMV-SPORT6.ccdb; Primer:
Onti; Site_2: EcoRV; Cloned unidirectionally. Primer:
Oligo dT. Average insert size 2.2 kb. Constructed by
Invitrogen. Note: this is a NCI_CGAP Library."
a 15 c 30 g 0 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:5331264"
/clone_lib="NCI_CGAP_Brn72"
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66.7%;
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Pred. No. 1.2e+06;
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                  M.D., Ph.D.
Brownstein (NHGRI), Shiraki
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BASE COUNT
ORIGIN
Search completed: March 13, 2003, 23:58:36 Job time: 2479 secs
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                                                                                                                                                                                           1352 AGCGGCGGGGGGGACCGCGGGGGGGGGGGGGGGAG 1386
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5276516"
/clone_ib="NIH_MGC_95"
/tissue_type="hippocampus"
/lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                 /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: Sall-xhoI (gtcgag); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library. "a 12 c 23 g 3 t
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